



FIGURE 1

FIGURE 41A

	₹1+ UV001-	
hUbiquitin Primary probe INVADER oligonucleotide ARRESTOR oligonucleotide FRET Probe Secondary target	5' -CGC CGA GAT CAC CTT TAC ATT TTC TAT CGT NH2-3' 5' -CCT TCC TTA TCC TGG ATC TTG GCA -3' 5'-ACG ATA GAA AAT GTA AAG GTG ATC-3' 5'-RED-CTC (Z28) TTC TCA GTG CG-3' 5'-CGC AGT GAG AAT GAG GTG ATC TCG GCG GT-3'	(SEQ ID NO:169) (SEQ ID NO:170) (SEQ ID NO:171) (SEQ ID NO:172) (SEQ ID NO:173)
m/r Ubiquitin, mouse (288C) Primary probe INVADER oligonucleotide 1 INVADER oligonucleotide 2 INVADER oligonucleotide 3 ARRESTOR oligonucleotide FRET Probe Secondary target	m/r Ubiquitin, mouse (288C, 516C, 744C, 972C), rat (247C, 475C, 703C, 931C) Primary probe S-CCG CCG AGA TCA CGG ATG TTG TAA TCA GAG A-NH2-3' INVADER oligonucleotide 2 S-GTG CAG GGT TGA CTC CTT CTC-3' S-GTG CAG GGT TGA CTC TTT CTC-3' S-GTG CAG CTC TTT C	(SEQ ID NO:174) (SEQ ID NO:175) (SEQ ID NO:176) (SEQ ID NO:177) (SEQ ID NO:177) (SEQ ID NO:172) (SEQ ID NO:173)
r/m GAPDH, rat (150C), mo Primary probe INVADER oligonucleotide ARRESTOR oligonucleotide FRET Probe Secondary target	mouse(166C) 5'-CGC CGA GAT CAC GTA GTT GAG GTC AAT GA-NH2-3' 5'-GAA TCA TAC TGG AAC ATG TAG ACC ATC-3' 5'-TCA TTG ACC TCA ACT ACG TGA TCT-3' 5'-RED-CTC (228) TTC TCA GTG CG-3' 5'-CGC AGT GAG AAT GAG GTG ATC TCG GCG GT-3'	(SEQ ID NO:179) (SEQ ID NO:180) (SEQ ID NO:181) (SEQ ID NO:172) (SEQ ID NO:173)
hGAPDH, 516C Primary probe INVADER oligonucleotide ARRESTOR oligonucleotide FRET Probe Secondary target	5'-CCG CCG AGA TCA CGA TGA TCT TGA GGC T-NH2-3' 5'-TGG TGC AGG AGG CAT TGC TC-3' 5'-CAG CCT CAA GAT TAC CGT GAT CT-3' 5'-RED-CTC (Z28) TTC TCA GTG CG-3' 5'-CGC AGT GAG AAT GAG GTG ATC TCG GCG GT-3'	(SEQ ID NO:182) (SEQ ID NO:183) (SEQ ID NO:184) (SEQ ID NO:172) (SEQ ID NO:173)

FIGURE 41B

hTGF-β		
Primary probe	5'- CCG TCA CGC CTC CTC GGC TC -3'	(SEQ ID NO:185)
INVADER oligonucleotide	5'- AGG CGA AAG CCC TCA ATT TCC CA-3'	(SEQ ID NO:186)
Stacker	5'-AAC CAC TGC CGC ACA-3'	(SEQ ID NO:187)
ARRESTOR oligonucleotide	5'-GAG CCG TGG AGG AGG CG-3'	(SEQ ID NO:188)
FRET Probe	5'-FL-CAC-(Z28)-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT -3'	(SEQ ID NO:190)
hMCP-1		
Primary probe	5'-CCG TCA CGC CTC CTT CGG AGT TTG GG NH2 -3"	(SEQ ID NO:191)
INVADER oligonucleotide	5' -GGG TTG TGG AGT GAG TGT TCA AGT A -3'	(SEQ ID NO:192)
Stacker	NO STACKER	
ARRESTOR oligonucleotide	5'-GGG-AAA-CTC-CGA-AGG- AGG-CG-3'	(SEQ ID NO:193)
FRET Probe	5'-FL-CAC-Z28-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT -3'	(SEQ ID NO:190)
hTNF-α		
Primary probe	5'-CCG TCA CGC CTC TCT GAC TGC CA NH2-3'	(SEQ ID NO:194)
INVADER oligonucleotide	5' -TTG TCA CTC GGG GTT CGA GAA GAT GAA-3'	(SEQ ID NO:195)
Stacker	5'-GGG CCA GAG GG-3'	(SEQ ID NO:196)
ARRESTOR oligonucleotide	5'-AGG CAG TCA GAG AGG CG-3'	(SEQ ID NO:197)
FRET Probe	5'-FL-CAC-Z28-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT -3'	(SEQ ID NO:190)
hIL-6		
Primary probe	5' -CCG TCA CGC CTC CTC ATT GAA TTNH2-3'	(SEQ ID NO:198)
INVADER oligonucleotide	5' -CCA AAA GTC CAG TGA TGA TTT TCA CCA GGC AAG TA -3'	(SEQ ID NO:199)
Stacker	5'-CAG ATT GGA AGC ATC CAT CT-3'	(SEQ ID NO:200)
ARRESTOR oligonucleotide	5'-GAT TCA ATG AGG AGG C-3'	(SEQ ID NO:201)
FRET Probe	5'-FL-CAC-(Z28)-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT -3'	(SEQ ID NO:190)

FIGURE 41C

hIL-1β		
Primary probe	5' -CCG TCA CGC CTC CAT CTG TTT AGG NH2-3'	(SEQ ID NO:202)
INVADER oligonucleotide	5' -CAG GTC CTG GAA GGA GCA CTT A-3'	(SEQ ID NO:203)
Stacker	5'-GCC ATC AGC TTC TTT GTT CTT GTÇ ATC-3'	(SEQ ID NO:204)
ARRESTOR oligonucleotide	5'-GCC CTA AAC AGA TGG AGG CG-3'	(SEQ ID NO:205)
FRET Probe	5'-FL-CAC-(Z28)-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:190)
hIL-2		
Primary probe	5'-CCG TCA CGC CTC CTC CAG TTG TAG NH2 -3'	(SEQ ID NO:206)
INVADER oligonucleotide	5'-AAA ATC ATC TGT AAA TCC AGC AGT AAA TGA -3'	(SEQ ID NO:207)
Stacker	5'-CTG TGT TTT CTT TGT AGA AC -3'	(SEQ ID NO:208)
ARRESTOR oligonucleotide	5' CTA CAA CTG GAG GAG GC -3'	(SEQ ID NO:209)
FRET Probe	5'-FL-CAC-(Z28)-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT- 3'	(SEQ ID NO:190)
hIL-8		
Primary probe	5'-CCG TCA CGC CTC CTC TCA GTT CT-NH2-3'	(SEO ID NO:210)
INVADER oligonucleotide	5'-GTG TGG TCC ACT CTC AAT CAA -3'	(SEQ ID NO:211)
Stacker	5'-TTG ATA AAT TTG GGG TGG AAA GGT TTG GA-3'	(SEQ ID NO:619)
ARRESTOR oligonucleotide	5'-AGA ACT GAG AGG AGG CG-3'	(SEQ ID NO:620)
FRET Probe	5'-FL-CAC-(Z28)-TGC TTC GTG G-3'	(SEQ ID NO:189)
Secondary target	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:190)
hIL-10		
Primary probe	5'-AAC GAG GCG CAC CAA ACT CAC TCA T-NH2-3'	(SEQ ID NO:621)
INVADER oligonucleotide	5'-GTC ATG TAG GCT TCT ATG TAG TTG ATG AAG ATG TA-3'	(SEQ ID NO:622)
Stacker	5'-GGC TTT GTA GAT GCC TTT CTC TTG GA-3'	(SEQ ID NO:623)
ARKES I OR Oligonucleotide FRET Prohe	5-AIG AGI GAG 111 GGI GCG-3'	(SEQ ID NO:624)
Secondary target	5-re-cac (229/-160 110 616 6-3 5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'	(SECTIONO:189)
		(070:010)

FIGURE 41D

hIL-4 Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe
hIL-4 Primary probe INVADER oligor Stacker ARRESTOR olig FRET Probe

5'-AAC GAG GCG CAC CTT CAA AAT GCC TAA-NH2-3' (SEQ ID NO:630) 5'-TGT CAC TCT CCT CTT TCC AAT TA-3'
5'-TGT CAC TCT CCT CTT TCC AAT TA-3'
5'-TTA GGC ATT TTG AAG GTG CGC-3'
5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'

FIGURE 41E

nCYP 1AZ, 1193G Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	5'-AAC GAG GCG CAC CGT TGT GTC CC-NH2-3' 5'- GGG AT G TAG AAG CCA TTC AGA-3' 5'-TTG TTG TGC TGT GGG GGA TG-3' 5'-GGG ACA CAA CGG TGC GC-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'	(SEQ ID NO:634) (SEQ ID NO:635) (SEQ ID NO:636) (SEQ ID NO:637) (SEQ ID NO:189) (SEQ ID NO:625)
hCYP 2B6, 343G Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	5'- CCG TCA CGC CTC CAC CAT ATC CC-NH2-3' 5'-CCA GCG GTT TCC ATT GGC AAA GAT CAA-3' 5'-CGG AAG AAT GGG TCG ACC ATG-3' 5'-GGG ATA TGG TGG AGG CG-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-FCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:638) (SEQ ID NO:639) (SEQ ID NO:640) (SEQ ID NO:189) (SEQ ID NO:190)
hCYP 2C19, 223G Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	5'-AAC GAG GCG CAC CGT TCC AGG C-NH2-3' 5'-CAT ATC CAT GCA GCA CCA TGA-3' 5'-CAA AAT ACA GAG TGA ACA CAG GGC C-3' 5'-GCC TGG AAC GGT GCG C-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'	(SEQ ID NO:642) (SEQ ID NO:643) (SEQ ID NO:644) (SEQ ID NO:645) (SEQ ID NO:625)
hCYP 2C9, 1554T Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	5'-CCG TCA CGC CTC ATG GAT AAT GCC C-NH2-3' 5'-CAG GTG AGA AAA GGC ATT ACA GAT AGT GAA AGC-3' 5'-CAG AGG AAA GAG AGC TGC AGG G-3' 5'-GGG CAT TAT CCA TGA GGC G-3' 5'-GGG CAT TAT CCA TGG GG G-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:646) (SEQ ID NO:647) (SEQ ID NO:648) (SEQ ID NO:49) (SEQ ID NO:189) (SEQ ID NO:190)

FIGURE 41F

5'-CCG TCA CGC CTC CCT GCT GAG AAA-NH2-3' 5'-CCC GAG GCA TGC ACG GCG GA-3' 5'-GGC AGG AAG GCC TCC-3' 5'-GGC AGG AGG GCG TCC-3' 5'-TTT CTC AGC AGG GAG GCG-3' 5'-TT CTC AGC AGG GAG GCG-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3' (SEQ ID NO:189)	A-NH2-3' C ATT C-3' (SEQ ID NO:656) (SEQ ID NO:657) (SEQ ID NO:189) SGT GAC GGT-3' (SEQ ID NO:190)	(SEQ ID NO:658) 3-3' 3-3' (SEQ ID NO:659) 3-3' (SEQ ID NO:660) 3-3' (SEQ ID NO:661) 3' (SEQ ID NO:189) 3' (SEQ ID NO:189)	AT ACC CC-NH2-3' (SEQ ID NO:662) AC-3' (SEQ ID NO:663) T AAC ATT C-3' (SEQ ID NO:664)
5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	5'-CCG TCA CGC CTC GCC CCA CA-NH2-3' 5'-CAG CAC AGG CTG TTG ACC ATC ATA AAA C-3' 5'-CTT TTC CAT ACT TTT TAT GAC ATT C-3' 5'-TGT GGG GCG AGG CG-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	5'-AAC GAG GCG CAC AGT TGA CCT TC-NH2-3' 5'-GTG ATG GCC AGC ACA GGG C-3' 5'-ATA CGT TCC CCA CAT TTT TC-3' 5'-TGA AGG TCA ACT GTG CGC-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'	5'-AAC GAG GCG CAC GTC ATA AAT ACC CC-NH2-3' 5'-GCC AGC ATA GGC TGT TGA CAC-3' 5'-AGA CTT TTC TAT ACT TTT TAT AAC ATT C-3' 5'-GGG GTA TTT ATG ACG TGC GC-3'
Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe	hCYP 3A4, 309C Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	hCYP 3A5 v2, 323T Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	hCYP 3A7, 231C Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide

FIGURE 41G

Primary probe INVADER oligonucleotide (h) INVADER oligonucleotide (r) Stacker ARRESTOR oligonucleotide FRET Probe	5'-CCG TCA CGC CTC CTG TCT GTG AT-NH2-3' 5'-TCC TGA CAG TGC TCA ATC AGG A-3' 5'-TCC TGA CAA TGC TCA ATG AGG A-3' 5'-GTC CCG GAT GTG GCC C-3' 5'-ATC ACA GAC AGG AGG CG-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:666) (SEQ ID NO:667) (SEQ ID NO:668) (SEQ ID NO:669) (SEQ ID NO:670) (SEQ ID NO:189) (SEQ ID NO:180)
h/rCYP 1A2 (813C/819C) Primary probe INVADER oligonucleotide (h) INVADER oligonucleotide (r) ARRESTOR oligonucleotide FRET Probe Secondary target	5'-AAC GAG GCG CAC GGA CTG TTT TCT GC-NH2-3' 5'-CTT GTC AAA GTC CTG ATA GTG CTC CTC-3' 5'-CTT GTT GAA GTC TTG ATA GTG TTC CTC-3' 5'-GCA GAA AAC AGT CCG TGC GC-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'	(SEQ ID NO:671) (SEQ ID NO:672) (SEQ ID NO:673) (SEQ ID NO:674) (SEQ ID NO:189) (SEQ ID NO:625)
rCYP 2B1, 1017T Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe	5'-CCG TCA CGC CTC ACT GCG GTC AT-NH2-3' 5'-GTG GAT AAC TGC ATC AGT GTA TGG CAT TTT C-3' 5'-CAA GGG TTG GTA GCC TGT GTG AGC C-3' 5'-ATG ACC GCA GTG AGG CG-3' 5'-ATG ACC (Z28) TGC TTC GTG G-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:675) (SEQ ID NO:676) (SEQ ID NO:677) (SEQ ID NO:678) (SEQ ID NO:189) (SEQ ID NO:190)
rCYP 2B2, 162T Primary probe INVADER oligonucleotide Stacker ARRESTOR oligonucleotide FRET Probe Secondary target	5'-CCG TCA CGC CTC AGA GCC AAT CAC-NH2-3' 5'-CGA TCA TCA AGG GAT GGT GGC CTG TGC-3' 5'-CTG ATC AAT CTC CTT TTG GAC TTT CTC TGC G-3' 5'-GTG ATT GGC TCT GAG GCG-3' 5'-GTG ATT GGC TCT GTG G-3' 5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'	(SEQ ID NO:679) (SEQ ID NO:680) (SEQ ID NO:681) (SEQ ID NO:189) (SEQ ID NO:189)

FIGURE 41H

5'-CCG TCA CGC CTC CTC TTC AAT TTC TG-NH2-3'	5'-CCC TGT CAA TTT CTT CAT GAA GTT TA-3'	5'-GGT ATT TCA TGA GGA TCA GGA GC-3"	5'-CAG AAA TTG AAG AGG AGG CG-3'	5'-FL-CAC (Z28) TGC TTC GTG G-3'	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT -3'
5-CCG T	5'-CCC T	5'-GGT A	5'-CAG	5'-FL-CA	5'-CCA G

5'-CCG TCA CGC CTC CTC GGC AGG-NH2-3'	5'-CCT GCC GAG GAG GCG-3'
5'-CAC AAT ATC GTA GGT AGG AGG TGC CTT AA-3'	5'-FL-CAC (Z28) TGC TTC GTG G-3'
5'-GCC CCA TCG ATC TCC TCC-3'	5'-CCA GGA AGC AAG TGG AGG CGT GAC GGT-3'

5'-AAC GAG GCG CAC TAG GCT TTG CT-NH2-3'	5'-TTC ATG TAG TCA GGG TCA TAG ACA ATT AAG A-3'	5'-TCC CCA GAA CCA TCG AGG AAA GG-3'	5'-AGC AAA GCC TAG TGC GC-3'	5'-FL-CAC (Z28) TGC TTC GTG G-3'	5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'

(SEQ ID NO:684) (SEQ ID NO:685) (SEQ ID NO:686) (SEQ ID NO:189) (SEQ ID NO:190)	(SEQ ID NO:687) (SEQ ID NO:688) (SEQ ID NO:689) (SEQ ID NO:690) (SEQ ID NO:189) (SEQ ID NO:625)

(SEQ ID NO:683)

EQ ID NO:6 EQ ID NO:6 EQ ID NO:6 EQ ID NO:6	(SEQ ID NO:190)
------------------------------------------------------	-----------------

(SEQ ID NO:695) (SEQ ID NO:696) (SEQ ID NO:697) (SEQ ID NO:698) (SEQ ID NO:698)	10.02
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FIGURE 411

5'-AAC GAG GAC AGA AGG CCC CTT-NH2-3'	5'-CCT TGA ACA GCA CCA GAA ATA GAC TGA GCA C-3' 5'-GGA AGA ACC CAG AGA CAC CAT CC-3'	5'-AAG GGG CCT TCT GTG CGC-3' 5'-FL-CAC (Z28) TGC TTC GTG G-3'	5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'		5'-AAC GAG GCG CAC GTT GTG ATA CCT T-NH2-3'	5-GAT GAA GGC CAT AAA TTA AAA TTG TGC-3'	5'-TGG GTA TGG AAC GTC C-3'	5'-AAG GTA TCA CAA CGT GCG C-3'	5'-FL-CAC (Z28) TGC TTC GTG G-3'	5'-CCA GGA AGC AAG TGG TGC GCC TCG TTT-3'
rCYP 4A2 Primary probe	INVADER oligonucleotide Stacker	ARRESTOR oligonucleotide FRET Probe	Secondary target	rCYP 4A3, 1235C	Primary probe	INVADER oligonucleotide	Stacker	ARRESTOR oligonucleotide	FRET Probe	Secondary target

(SEQ ID NO:699) (SEQ ID NO:700) (SEQ ID NO:701) (SEQ ID NO:189) (SEQ ID NO:625) (SEQ ID NO:703) (SEQ ID NO:704) (SEQ ID NO:705) (SEQ ID NO:706) (SEQ ID NO:189) (SEQ ID NO:625)

FIGURE 47A

Oligo sequence descriptions: 5' to 3' direction, 2'-Ome nts are bolded and underlined, internal modifications defined in ()

Oligo Type	Oligo Sequence (5' to 3')	Modification	SEQ ID NO
hTNF-a			
probe	ccg ccg aga tca ctc tga ctg cct NH2	3' Amine	002
invader	ttg tca ctc ggg gtt cga gaa gat gaa		710
stacker	ggg cca gag ggc tga tta g	all 2'Ome bases	711
stacker	ggg cca gag ggc tga tta	all 2'Ome bases	712
stacker	ggg cca gag ggc tg at	all 2'Ome bases	713
stacker	ggg cca gag ggc t	all 2'Ome bases	714
stacker	<u> </u>	all 2'Ome bases	715
arrestor	agg cag tea gag tga te	all 2'Ome bases	716
arrestor	agg cag tea gag tga tet c	all 2'Ome bases	717
SRT	cggaagaagcagttggtgatctcggcggNH2	3' Amine	718
FRET probe	Fcaac(Cy3)gcttcctccg		719
probe	ccg tca cgc ctc tct gac tgc ct NH2	3' Amine	720
invader	ttg tca ctc ggg gtt cga gaa gat gaa)	731
stacker	ggg cca gag ggc tga tta g	all 2'Ome bases	12)
arrestor	agg cag tea gag agg eg	all 2'Ome bases	723
SRT	cggaagaagcagttggaggcgtgacggtNH2	3'base 2'Ome. 3'Amine	727
FRET probe	Fcaac(Cy3)gcttcctccg		725
probe	oca toa cac oto tot aac tac ota aNH2		;
invader	tto toa ctc aga att caa aaa aat aaa	שׁבְּבְּבָּבְּבָּבְּבְּבְּבָּבְּבְּבְּבְּבְּבְּבְּבְּבְּבְּבְּבְּבְּ	726
arrestor	cca ddc adt cad ada ddc d	2004 cmO'c 11c	171
SRT	cqqaaqaaqcaqttqqaqqcqtoacqqtNH2	3'hose 2'Amino	728
FRET probe	Fcaac(Cy3)gcttcctccg		730
probe	cca cca aga tca ctc tga cto cc NH2	2. Amino	
invader	ttg tca ctc ggg gtt cga gaa gat gaa	<u> </u>	727
stacker	tog gcc aga ggg ctg att a	all 2'Ome bases	732
arrestor	agg cag tea gag tga te	all 2'Ome bases	737
SRT	cggaagaagcagttggtgatctcggcggNH2	3' Amine	735
FRET probe	Fcaac(Cy3)gcttcctccg		736
probe	ccg ccg aga tca ctg atc tga ctg NH2	3' Amine	7.27
invader	ctt gtc act cgg ggt tcg aga aga c)	738

FIGURE 47B

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FIGURE 47C

FIGURE 47D

805 806 807 808 809	810 812 813 814 815 816	817 818 819 820 821	823 824 825 827 828 830	831 832 833 834 835
3' Amine 3' Amine <u>all 2'Ome bases,3' Amine</u> 3' 2 last base <u>2' Ome</u> , 3' Amine	3' Amine all 2'Ome bases all 2'Ome bases 3'base 2'Ome, 3'Amine	3' Amine all 2'Ome bases all 2'Ome bases 3'base 2'Ome, 3'Amine	3' Amine 3' Amine 3' Amine all 2'Ome bases all 2'Ome bases 3' 3bases 2'Ome	3' Amine 3' Amine <u>all 2'Ome bases</u> 3'base 2'Ome , 3'Amine
ccg ccg aga tca ctc tcc ttg aat cct NH2 ccg ccg aga tca ctc tcc tca ttg aat ccNH2 cca aaa gtc cag tga tga tt tca cca ggc aag a agg att caa tga gga agg atc tNH2 cggaggaagcagttggtgatctcggcggNH2 Fcaac(CA)nctfcctcc	ccg tca cgc ctc ctc att gaaNH2 cca gtg atg att ttc acc agg caa gta tcc aga ttg gaa gca tcc atc t ttc aat gaa gag ga gc cggaagaagcagttggaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	ccg tca cgc ctc ctc att gaNH2 cca gtg atg att ttc acc agg caa gta atc cag att gga agc atc cat ct ttc aat gag gag gc cggaagaagcagttggaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	ccg tca cgc ctc ctc att gaa tgNH2 ccg tca cgc ctc ctc att gaa taNH2 ccg tca cgc ctc ctc att gaa taNH2 ccg tca cgc ctc ctc att gaa ttNH2 cca aaa gtc cag tga tga ttt tca cca ggc aag ta cagattggaagcatccatct gattcaatgaggaggcg ccaggaagcaagtggaggc ccaggaagcaagtggaggcgtgacggu Fcac(Z21)tgcttcgtgg	ccg tca cgc ctc ctt cgg agt ttg gtNH2 ccg tca cgc ctc ctt cgg agt ttg gtt NH2 ggg ttg tgg agt gag tgt tca agt a aac cca aac tcc gaa ggc ggc gtg NH2. cggaagaagcagttggaaggcgtgacggtNH2.
hlL-6 probe probe invader arrestor SRT FRET probe	probe invader stacker arrestor SRT FRET probe	probe invader stacker arrestor SRT FRET probe	probe probe probe invader stacker arrestor SRT FRET probe	hMCP-1 probe probe invader arrestor SRT

FIGURE 47E

FRET probe	Fcaac(Cy3)gcttcctccg		836
probe probe invader arrestor SRT FRET probe	gcc gtc acg cct ctt tgg gtt tgc ttg tc NH2 gcc gtc acg cct ctt tgg gtt tgc ttg tNH2 tggagtgagtgttcaagtcttcggaga gacaagcaaaccaaagagcg cgaagaagaagcagttggaggcgtgacggcNH2 Fcaac(Cy3)gcttcctccg	3' Amine 3' Amine all 2'Ome bases 3'2 bases 2'Ome , 3'Amine	837 838 839 840 841
probe probe invader arrestor SRT SRT SRT SRT SRT SRT SRT SRT SRT	cct gtc tcg ctg cct tcg gag ttt ggg cct gtc tcg ctg cct tcg gag ttt gg ggg ttg tgg agt gag ttt a ga ggg ttg tgg agt gag tt ca agt a ccc.aaa ctc.cga agg cag.cg cggaggaagcagttggcagcgagaacaggNH2 cggaggaagcagttggcagcg(Amino dA)ggNH2 cggaggaagcagttggcagcg(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gacagac(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gcgagac(Amino dA)gacaggNH2 cggaggaagcagttggc(Amino dA)gcg(Amino dA)gacaggNH2 Fcaac(Cy3)gcttcctccg	all 2'Ome bases 3' last base 2'Ome, 3' Amine Amino dA modification	843 844 845 846 847 850 851 853
probe invader arrestor SRT FRET probe	gcc gtc acg cct ctg gga cac ttg ctg cNH2 gcc aca atg gtc ttg aag atc aca gct tct ta gca gca agt gtc cca gag gcg NH2 cggaagaagcagttggaggcgtgacggcNH2 Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases.3' Amine 3'2 bases 2'Ome , 3'Amine	855 856 857 858 859
probe invader arrestor SRT FRET probe	ccg tca cgc ctc ctt cgg agt ttg gg NH2 99g ttg tgg agt gag tgt tca agt a 5'-ggg-aaa-ctc-cga-agg-agg-cg-3' ccaggaagcaagtggaggcgtgacggu Fcac(Z21)tgcttcgtgg	3' Amine all 2'Ome bases 3' 3bases 2'Ome	860 861 862 863
probe invader arrestor SRT FRET probe	cgc cga gat cac ctt cgg agt ttg ggNH2 ggg ttg tgg agt gat tca agt a ccc aaa.ctc cga agg tga tc cggaagaagcagttggtgatctcggcggNH2 Fcaac(C)3)gcttcctccg	3' Amine all 2'Ome bases 3' Amine	865 866 867 868 869

FIGURE 47F

probe invader	aac gag gcg cac ctt cgg agt ttg gg NH2	3' Amine	870
arrestor SRT FRET probe	999 '19 '19 '19 '19 '19 '10 '10 '10 '10 '10 '10 '10 '10 '10 '10	all 2'Ome bases 3' last 5 bases 2'Ome , 3' Amine	871 872 873 874
probe	ccg tca cgc ctc ctt cgg agt ttg g NH2	3' Amine	875
stacker	999 iig igg agi gag iga ica agi a git igc itg icc agg igg	all 2'Ome bases	876 877
arrestor	cca aac tee gaa gga gge g	all 2'Ome bases	878
SRT FRET probe	cggaagaagcagttggaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	3'base 2'Ome , 3'Amine	879 880
probe	ccg tca cgc ctc ctt cgg agt ttg NH2	3' Amine	881
invader	ggg ttg tgg agt gag tgt tca agt a		882
stacker	att ttg ctt gtc cag gtg g	all 2'Ome bases	883
arrestor	cca aac tcc gaa gga ggc g	all 2'Ome bases	884
SRT EDET probe	cggaagaagcagttggaggcgtgacggtNH2	3'base 2'Ome , 3'Amine	885
Pione	rcaac(cys)gciicciccg		886
probe	ccg tca cgc ctc ctt cgg agt ttNH2	3' Amine	887
invader	ggg ttg tgg agt gag tgt tca agt a		888
stacker	agg ttt get tat een gat g	all 2'Ome bases	889
arrestor	cca aac tcc gaa gga ggc g	all 2'Ome bases	890
RI I	cggaagaagcagttggaggcgtgacggtNH2	3'base 2'Ome , 3'Amine	891
FRET probe	Fcaac(Cy3)gcttcctccg		892
probe	ccgtcacgcctccggagtttgggNH2	3' Amine	893
invader	gtt gtg gag tga gtg ttc aag tat ta		894
stacker	ttt get tgt eca ggt ggt eca g	all 2'Ome bases	895
arrestor	ccc aga ctc cgg agg cg	all 2'Ome bases	896
SRT	cggaagaagcagttggaggcgtgacggtNH2	3'base 2'Ome , 3'Amine	897
FRET probe	Fcaac(Cy3)gcttcctccg		868
probe	cgc cga gat cac cgg agt ttg ggNH2	3' Amine	899
nvader	gtt gtg gag tga gtg ttc aag tat ta		006
stacker	ttt get tgt eca ggt ggt eca g	all 2'Ome bases	901
arrestor	cta gtg gcc tca aac cc	all 2'Ome bases	902
SRT	cggaagaagcagttggtgatctcggcggNH2	3' Amine	903
FRET probe	Fcaac(Cy3)gcttcctccg		904

FIGURE 47G

hUbiquitin probe probe invader arrestor SRT FRET probe	cgc cga gat cac ctt tac att ttc tat cgt cgc cga gat cac ctt tac att ttc tat cgt NH2 5' –cct tcc tta tcc ttg atc ttg gca -3' acg ata gaa aat gta aag gtg atc. 5'-cgc agt gag aat gag gtg atc tcg gcggt-3' 5'-cgc agt gag gtg gtc tcg gcggt-3' 5'-Red-ctc-Z21-ttc tca gtg cg-3'	3' Amine all 2'Ome bases 3' last 3 bases 2'Ome.	905 906 907 908 909
hIL-2 probe invader stacker arrestor SRT FRET probe	gtttcttttgtgtctccgcactgccNH2 cca gca gta aat gct cca gtt gta ga tag aac ttg aag tag gtg c. caa aga aaa cac agg agg c. ccaggaagcaagtggaggcgtgacg Fcac(Z21)tgcttcgtgg	3' Amine all 2'Ome bases all 2'Ome bases 3' 3bases 2'Ome	0 0 0 0 0 0 1 0 0 0 0 0 0
probe invader stacker arrestor SRT FRET probe	aac gag gcg cac ctg tgt ttt ctt tg NH2 cca gca gta aat gct cca gtt gta ga tag aac ttg aag tag gtg c. caa aga aga aga tgc c caa aga aga aga tgc g ccaggaagcaagtgggcgcctcgttt Fcac(Z21)tgcttcgttg	3' Amine all 2'Ome bases all 2'Ome bases 3' last 3 bases 2'Ome	917 918 919 920 921
probe invader stacker arrestor SRT FRET probe	ccg tca cgc ctc ctc cag ttg tag NH2 aaa atc atc tgt aaa tcc agt aaa tga ctg tgt ttt ctt tgt aga ac cta caa ctg aag gag gc ccaggaagcaagtggaggcgtgacggu Fcac(Z21)tgcttcgtgg	3' Amine 5' 6 bases 2'Ome all 2'Ome bases all 2'Ome bases 3' 3bases 2'Ome	923 924 925 926 927
probe invader stacker arrestor SRT FRET probe	aac gag gcg cac ctc cag ttg tag NH2 aaa atc atc tgt aaa tcc agc agt aaa tga ctg tgt ttt ctt tgt aga ac. cta caa ctg gag gtg cg. ccaggaagcaagtggtgcgcctcgttt Fcac(Z21)tgcttcgtgg	3' Amine 5' 6 bases 2'Ome all 2'Ome bases all 2'Ome bases 3' last 3 bases <u>2'Ome</u>	929 930 931 932 933

FIGURE 47H

935 936 937 939 940	941 942 943 945 946	947 948 949 950 952 953	955 956 957 958 959	960 961 962 963 964	996 968 969
3' Amine all 2'Ome bases all 2'Ome bases, 3' amine 3' 3bases 2'Ome	3' Amine all 2'Ome bases all 2'Ome bases 3' last 3 bases 2'Ome	3' Amine 3' Amine 3' Amine 5' 6 bases 2'Ome all 2'Ome bases all 2'Ome bases 3' 3bases 2'Ome	3' Amine all 2'Ome bases,3' Amine 3'2 bases 2'Ome , 3'Amine	3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine 3'2 bases 2'Ome, 3'Amine	All 2' Ome
ccg tca cgc ctc ctg tgt ttt ctt tgt aNH2 gta aat cca gca gta aat gct cca gtt gta ga gaa_ctt_gaa_gta_gt gca_ctg_tt tacaaagaaaacacaggaggcgtNH2 ccaggaagcaaqtggggggtgacggu Fcac(Z21)tgcttcgtgg	aac gag gcg cac ctg tgt ttt ctt tgt aNH2 gta aat cca gca gta aat gct cca gtt gta ga gaa ctt gaa gta ggt gca ctg tt tac aaa gaa aac aca ggt gcg ccaggaagcaagtggtgcgcctcgttt Fcac(Z21)tgcttcgtgg	ccg tca cgc ctc ctc cag ttg taa NH2 ccg tca cgc ctc ctc cag ttg tat NH2 ccg tca cgc ctc ctc cag ttg tac NH2 aaa atc atc tgt aaa tcc agc agt aaa tga ctg tgt ttt ctt tgt aga ac cta caa ctg gag gag gc ccaggaagcaagtggaggggggacggu	gcc gtc acg cct ccc ttc ttg atg NH2 ttc tag aca ctg aag atg ttt cag ttc tgt gga cat gcc caa gaa ggg agg cg NH2 cggaagaagcagttggaggcgtgacggcNH2 Fcaac(Cy3)gcttcctccg	ccg tca cgc ctc taa ttc cat tca aaa tca tct NH2 cat cct ggt gag ttt ggg att ctt gta att tat a gta aat cca gca gta aat gct cca gNH2 aga tgc taa ttt tga atg gaa tta gag gcg NH2 cggaagaagcagttggaggcgtgacgcNH2 cggaagaagcagttggaggcgtgacgcNH2 Fcaac(Cy3)gcttcctccg	ccg ccg aga tca cct gtg ttt tct ttg ta gta aat cca gca gta aat gct cca gtt gta ga gaa ctt gaa gta ggt gca ctg tt gaa ctt gaa gta ggt gca ctg tt
probe invader stacker arrestor SRT FRET probe	probe invader stacker arrestor SRT FRET probe	probe probe probe invader stacker arrestor SRT FRET probe	probe invader arrestor SRT FRET probe	probe invader stacker arrestor SRT FRET probe	probe invader stacker stacker

FIGURE 47I

stacker stacker arrestor SRT FRET probe	gaa ctt gaa gta ggt gca ctg tt gaa ctt.gaa gta ggt gca ctg tt tac aaa gaa aac aca ggt gat ct cggaggaagcagttggtgatctcggcggNH2 Fcaac(Cy3)gcttcctccg	5' 3bases 2'Ome 5' 6bases 2'Ome All 2' Ome 3' 2 last base <u>2' Ome</u> , 3' Amine	970 971 972 973
probe invader arrestor SRT FRET probe	aac gag gcg cac cct tct tgg gca tgNH2 ttc tag aca ctg aag atg ttt cag ttc tgt gga cat gcc caa gaa ggg tgc gNH2 cggaagaagcagttggtgccctc gttaa NH2 Fcaac(Cy3)gcttcctccg	3' Amine <u>all 2'Ome bases</u> 3' last 5 bases <u>2'Ome,</u> 3' Amine	975 976 977 978 979
probe invader stacker arrestor SRT FRET probe	aac gag gcg cac taa ttc cat tca aaa tca tct cat cct ggt gag ttt ggg att ctt gta att tat a gta aat cca gca gta aat gct cca gNH2 aga tta att gaa tta gt NH2 aga tta att gaa tta gt NH2 cggaagaagcagttggtgcctcgttaaNH2 Fcaac(Cy3)gcttcctccg	all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine 3' last 5 bases 2'Ome, 3' Amine	980 981 982 983 985
hIL-4 probe invader invader arrestor arrestor arrestor SRT FRET probe	cct gtc tcg ctg cca gtt gtg ttc ttg gag NH2 ccc tgc aga agg ttt cct tct a ccc tgc aga tgg ttt cct tct a ctc caa gaa cac aac tgg cag cNH2 ctc caa gaa cac aac tgg cag cga NH2 ctc caa gaa cac aac tgg cag cga NH2 ctc caa gaa cac aac tgg cag cga NH2 ctc caa gaa cac aac tgg cag cga NH2 ctc caa gaa cac aac tgg cag cga NH2 cfc caa gaa cac aac tgg cag cga NH2 cggagggaagcagttggcagcgagacaggNH2 Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine 3' last base 2'Ome, 3' Amine	986 987 999 992 992
probe probe invader arrestor SRT FRET probe	aac gag gcg cac ctt gga ggc agc aaa NH2 aac gag gcg cac ctt gga ggc agc aaNH2 aag gtt tcc ttc tca gtt gtg tta ctt tgc tgc ctc caa ggt gcg NH2 cggaggaagcagttggtgcgcctcgttaa NH2 Fcaac(Cy3)gcttcctccg	3' Amine 3' Amine all 2'Ome bases.3' Amine 3' last 5 bases 2'Ome, 3' Amine	994 995 997 998 999
probe invader arrestor	cag tca cgt ctc tgg agg cag caa aga tg NH2 aag gtt tcc ttc tca gtt gtg ttc ta cat ctt tgc tgc ctc cag aga cg NH2	3' Amine all 2'Ome bases,3' Amine	1000 1001 1002

FIGURE 47J

SRT FRET probe	gctactgagatgaaggagacgtgactgtaNH2 Fcttc(Cy3)tctcagtagc	3' Amine	1003 1004
probe invader arrestor SRT FRET probe	aac gag gcg cac ctt gga ggc agc aaa g NH2 aag gtt tcc ttc tca gtt gtg tta ctt tgc tgc ctc caa ggt gcg NH2 cggaggaagcagttggtgcgcctcgttaa Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases,3' Amine 3' last 5 bases 2'Ome	1005 1006 1007 1008
mIL-2 probe invader arrestor SRT FRET probe	cgc cga gat cac ccc ttt agt ttt aca aca gtNH2 gaa ttg gca ctc aaa tgt gtt gtc aga ga act gtt gta aaa cta aag ggg gtg atc t NH2 cggaggaagcggttggtgatctcgg cg NH2 Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases,3' Amine 3' last two bases are 2' Ome , 3' Amine	010 1011 1013 1013
probe invader arrestor arrestor arrestor SRT FRET probe	tgc cgc cga gat cac ccc ttt agt ttt aca aca gtNH2 gaa ttg gca ctc aaa tgt gtt gtc aga ga act gtt gta aaa cta aag ggg gtg NH2 act gtt gta aaa cta aag ggg gtg at NH2 act gtt gta aaa cta aag ggg gtg at ctNH2 act gtt gta aaa cta aag ggg gtg at ctNH2 act gtt gta aaa cta aag ggg gtg at ctNH2 cggaggaagcggttggtgatctcggcggcaNH2 Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine 3' Last 2bases 2'Ome, 3' Amine	1015 1016 1017 1018 1020 1021
probe probe invader arrestor SRT FRET probe	gc cgc cga gat cac ccc ttt agt ttt aca aca gtNH2 c cgc cga gat cac ccc ttt agt ttt aca aca gtNH2 gaa ttg gca ctc aaa tgt gtt gtc aga ga act gtt gta aaa cta aag ggg gtg at NH2 cggaggaagcggttggtgatctcggcggcaNH2 cgaac(Cy3)gcttcctccg	3' Amine 3' Amine <u>all 2'Ome bases.3' Amine</u> 3' Last 2bases 2'Ome, 3' Amine	1023 1024 1025 1026 1027
probe invader arrestor SRT FRET probe	aác gag gcg cac ccc ttt agt ttt aca aca gt NH2 gaa ttg gca ctc aaa tgt gtt gtc aga ga agtaactgttgtaaaactaaaggggtgcg cggaggaagcagttggtgcgcctcgttaa Fcaac(Cy3)gcttcctccg	3' Amine all 2'Ome bases.3' Amine 3' last 5 bases 2'Ome	1029 1030 1031 1032
probe	aac gag gcg cac ccc ttt agt ttt aca aca gt NH2	3' Amine	1034

FIGURE 47K

invader arrestor SRT FRET probe	gaa ttg gca ctc aaa tgt gtt gtc aga ga agt aac tgt tgt aaa act aaa ggg gtg cg NH2 cggaggaagcagttggtgcgcctcgttaa Fcaac(Cy3)gcttcctccg	all 2'Ome bases,3' Amine 3' last 5 bases 2'Ome	1035 1036 1037 1038
probe invader	cogtoacgcotccctttagttttacaacNH2 gaa ttg goa ctc aaa tot gtt gtc aga ga	3' Amine	1039
stacker arrestor SRT	agt tac tot gat att got gat gaa att oto ag gttgtaaaactaaaggggaggcg cggaagaagcagttggaggcgtgacggtNH2	all 2'Ome bases, all 2'Ome bases, 3'base 2'Ome, 3'Amine	1040 1041 1043
FRET probe	Fcaac(Cy3)gcttcctccg		1044
probe invader	cgccgagatcaccctttagtttacaacNH2 gaa ttg gca ctc aaa tot gtt gtc aga ga	3' Amine	1045
stacker	agt tac fet gat aft get gat gaa aft ete ag	All 2:Ome	1047
SRT FRET probe	<u>รูกษาลสสสสเศสสสูษูษูษูตุลเต</u> cggaagaagcagttggtgatctcggcggNH2 Fcaac(Cv3)ncttcricco	All Z'Ome 3' Amine	1048
	BoomonoB/o (o) boomo		1050
probe invader	cogtcacgcctcccctttagttttacaaNH2 qaa tto qoa ctc aaa tot ott otc aga ga	3' Amine	1051
stacker	cagttactctgatattgctgatgaaattctca	All 2'Ome	1053
arrestor	gitgtaaaactaaagggagggcg	All 2'Ome	1054
FRET probe	csgaagaagkaguggaggggggggggggggggggggggggg	3 base 2.0me , 3 Amine	1055 1056
probe	ccgtcacgcctcccctttagttttacaaNH2	3' Amine	1057
stacker	gaz ing gaz ara igi gir gir gir aga ga cagitacicigatattgctgatgaaattctca	All 2'Ome	1058
arrestor	gitgtaaaactaaaggggaggcg	All 2'Ome	1060
FRET probe	ccaggaagcagttggaggcgtgacgg tNH2 Fcaac(Cy3)gcttcgtgg	3' 2 bases 2'Ome , 3'Amine	1061 1062
mIL-10			
probe	ccg tca cgc ctc ccg tta gct aag at NH2	3' Amine	1063
invader stacker	cga ggt ttt cca agg agt tgt tta ccc tgg atc aga ttt aga gag c	2.0 m O: 1 =	1064
arrestor SRT	atc tta gct aac ggg agg cg cggaagaagcagttggaggcgtgacggtNH2	all 2'Ome bases. 3'base 2'Ome, 3'Amine	1065 1067

	FIGURE 47L	7	"Replacement Sheet"
FRET probe	Fcaac(Cy3)gcttcctccg		1068
probe invader	ccg toa cgc ctc agt tgt ttc cgt tNH2 aga ggt aca aac gag att ttc caa ggc	3' Amine	1069
stacker arrestor		all 2'Ome bases,	1070
SRT FRET probe	ccaggaagcaagtggaggcgtgac ggu Fcac(Z21)tgcttcgtgg	all & Vine pases, 3' 3bases 2'Ome	10/2 1073 1074
probe	ccg tca cgc ctc ccg tta gct aNH2	3' Amine	1075
stacker	caa avy ayy iii too aay gag iig a aga too ctq gat cad att tag aga got c	all 2'Ome hases	1076
arrestor SPT	tag cta acg gaa aga ggc g	all 2'Ome bases.	1078
FRET probe	v.aggaagvaagtgaagvgtgac ggu Fcac(Z21)tgcttcgtgg	3. 3bases 2'Ome	1079 1080
probe	ccg tca cgc ctc ccg tta gNH2	3' Amine	1081
invader	aga ggt aca aac gag gtt ttc caa gga ga		1082
stacker arrestor	cia aga icc cig gal cag att tag aga g ctaacogaaacaagaggc	All 2'Ome	1083
SRT	ccaggaagcaagtggaggcgtgacgg <u>u</u>	3' 3bases 2'Ome	1084
FRET probe	Fcac(Z21)tgcttcgtgg		1086
hIFN-√			
probe	aac gag gcg cac ctt acc aat gcc taa gaa aag agt tNH2	3' Amine	1087
invader	tgc att att ttt ctg tca ctc tcc tct ttc caa tta		1088
arrestor	age tet tit ett agg eat tit gaa ggt geg NH2	all 2'Ome bases, 3' Amine	1089
FRET probe	cggaggaagcagugggcccc <mark>gmaa</mark> nnz Fcaac(Cy3)gcttcctccg	3. last 5 bases 2.0me	1090 1091
probe	cag toa cgt ctc tct tca aaa tgc cta aga aaa gag tNH2	3' Amine	1092
arrestor	ict gea ita itt itc tgt cac fot eet ett tee aat a act ett tte tta one att tto aan ana nac nNH2		1093
SRT	gctactgagatgaaggagacgtgactgtaNH2	all 2'Ome bases,3' Amine	1095
FRET probe	Fcttc(Cy3)tctcagtagc		1096
miFN -γ probe	aac gag gcg cac cct ttt gcc agt tcc NH2	3' Amine	1097

FIGURE 47M

invader arrestor SRT FRET probe	gct ctg cag gat ttt cat gtc acc ata gag gaa ctg gca aaa ggg tgc gNH2 gctactgagatgaaggagacgtgactgtaNH2 Fcttc(Cy3)tctcagtagc	all 2'Ome bases.3' Amine all 2'Ome bases.3' Amine	1098 1099 1100
probe invader	aac gag gcg cac cct ttt gcc agt NH2 gct ctg cag gat ttt cat gtc acc ata	3' Amine	1102
stacker arrestor	tee tee aga tat eea aga aga gae te act ede aga ago eeg ee	all 2'Ome bases	104
SRT	cgg agg aaag cag ttg gtg cgc ctc guu aa NH2	3' last 5 bases 2'Ome	1106
SRT	cgg aag aaag cag ttg gtg cgc ctc guu aa NH2	3' last 5 bases 2'Ome	1107
rkei probe	Fcaac(C)3)gcttcctccg		1108
probe	gcc gca cgc ctt ttg cca gt NH2	3' Amine	1109
invader	got otg cag gat tit cat gtc acc ata		1110
arrestor	act not assessed for aga aga gac ic	all 2'Ome bases	1111
SRT	cgg agg aag cag ttg cgg cgt gcg gca NH2	all & Ollie pases	1112
FRET probe	Fcaac(Cy3)gcttcctccg		1114
probe	aac gag gcg cac cct ttt gcc agt tc NH2	3' Amine	1115
stacker	got oug cag gai iii cat gic acc ata cto cag ata too aag aag aga cto		1116
arrestor	gaa ctg gca aaa ggg toc g	all 2'Ome bases	111/
SRT	cggaggaagcagttggtgcgcctcgttaaNH2	3' Jasts bases 2'Ome	0 7
FRET probe	Fcaac(Cy3)gcttcctccg	14450 Dates	1120
hlt-8			
probe	cca toa cac ctc ctt aac aaa act aca coNH2	3' Amine	7
probe	ccg tca cgc ctc ctt ggc aaa act gca cca NH2	3' Amine	1121
invader	ctt tat gca ctg aca tct aag ttc ttt agc act ca		1123
arrestor	tgg tgc agt ttt gcc aag gag gcg NH2	all 2'Ome bases.3' Amine	1124
arrestor	tog toc agt ttt gcc aag gag gcg tg NH2	all 2'Ome bases, 3' Amine	1125
SKI FRET probe	cggaagaagcagttggaggcgtgacg gc NH2 Fcaac(Cy3)gcttcctccg	3'2 bases 2'Ome , 3'Amine	1126
			171
probe	ccg tea cgc ctc cat ctt cac tga ttc ttg gNH2	3' Amine	1128
invader	agt att gaa ata aat tto ctt gaa att tea eto ga	3. Amine	1129
			081

FIGURE 47N

stacker	gat acc aca gag aat gaa tttt	all 2'Ome bases	1131
arrestor	tcc and ant can the aga the agg co NH2	all 2'Ome bases, 3' Amine	1132
arrestor	tcc aag aat cag tga aga tgg agg cgt gNH2	all 2'Ome bases, 3' Amine	1133
arrestor	g aat cag tga aga tgg agg cg	all 2'Ome bases	1134
SRT	cggaagaagcagttggagggggggggcNH2	3'2 bases 2'Ome , 3'Amine	1135
i NET prope	r caac(U)3)gciicciccg		1136
probe	ccg tca cgc cct tgg ctc aat ttt gct NH2	3' Amine	1137
invader	cca ttc aat tcc tga aat taa agt tcg gat att ctc ttg gca		1138
invader	cc tga aat taa agt tcg gat att ctc ttg gca	5' 10 bases are 2'Ome	1139
invader	cc tga aat taa agt tcg gat att ctc ttg gca		1140
arrestor	age asa att gag eca agg gag geg NH2	all 2'Ome bases, 3' Amine	1141
arrestor sip.r	age aga att gag eeg agg gag tgNH2	all 2'Ome bases, 3' Amine	1142
FRET probe	oggaagaagaaguggagguggaaggaagaagCoggaagaagCoggaagaagaagaagaagaagaagaagaagaagaagaaga	3.2 bases Z.Ume , 3.Amine	1143
			44.
probe	ccg tca cgc ctc cat ctt cac tga ttc ttg NH2	3' Amine	1145
invader	tto tag caa acc cat toa att cot gaa att aaa gtt ogg ata tto ta		1146
invader	cc cat tca att cct gaa att aaa gtt cgg ata ttc ta	5' 10 bases 2'Ome	1147
invader	cc cat tca att cct gaa att aaa gtt cgg ata ttc ta		1148
arrestor	cca agg gcc aag gag gcg tNH2		1149
ייייי דותה	cggaagaagcagttggaggcgtgacg gc NH2	3'2 bases 2'Ome , 3'Amine	1150
rke: probe	Fcaac(Cy3)gcttcctccg		1151
probe	ccg tca cgc ctc cat ctt cac tga ttc NH2	3' Amine	1152
invader	agt gtt gaa gta gat ttg ctt gaa gtt tca ctg ga		1153
stacker	ttg gat acc aca gag aat gaa tt	all 2'Ome bases	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
SRT	cggaagaagcagttggaggcgtgacgtNH2	3'hase 2'Ome 3'Amine	
FRET probe	Fcaac(Cy3)gcttcctccg		1156
probe	ccg tca cgc ctc cat ctt cac tga tt NH2	3' Amine	1157
invader	agt gtt gaa gta gat ttg ctt gaa gtt tca ctg ga		1158
stacker	ctt gga tac cac aga gaa tga att		1159
SRT	cggaagaagcagttggaggcgtgacggtNH2	3'base 2'Ome , 3'Amine	1160
FRET probe	Fcaac(Cy3)gcttcctccg		1161
probe	ccg toa cgc ctc cat ctt cac tga ttc ttg NH2	· 3' Amine	1162
invader	agt gtt gaa gta gat ttg ctt gaa gtt tca ctg ga		1163
helper	ata-cca-cag-aga-atg-aat-ttt-ttt-atg	all 2'Ome bases	1164
arrestor	tec aag aat cag tga aga tgg agg cgt gNH2	all 2'Ome bases, 3' Amine	1165

FIGURE 470

SRT FRET probe	cggaagaagcagttggaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	3'base 2'Ome , 3'Amine	1166 1167
SRT FRET probe	cggaagaagcagttggtgatctcggcggNH2 Fcaac(Cy3)gcttcctccg	3' Amine	1168 1169
SRT FRET probe	cggaagaagcagttggaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	3'base 2'Ome , 3'Amine	1170
SRT FRET probe	ccaggaagcaagtggaggcgtgac ggu Fcac(Z21)tgcttcgtgg	3' 3bases 2'Ome	1172
SRT FRET probe	cggaggaagcagttggtgatctcggc ggN H2 Fcaac(Cy3)gcttcctccg	3' 2 last base <u>2' Ome</u> , 3' Amine	1174
SRT FRET probe	cggaagaagcagttggaggcgtgacg gc NH2 Fcaac(Cy3)gcttcctccg	3'2 bases 2'Ome , 3'Amine	1176
SRT FRET probe	ccaggaagcaagtggtgcgcctcgttt Fcac(Z21)tgcttcgtgg	3' last 3 bases 2'Ome	1178
SRT FRET probe	cggaggaagcagttggtgcgcctc gttaaNH2 Fcaac(Cy3)gcttcctccg	3' last5 bases 2'Ome	1180
SRT FRET probe	cggaggaagcggttggtgatctcggcgg <u>ca</u> NH2 Fcaac(Cy3)gcttcctccg	3' Last 2bases 2'Ome, 3' Amine	1182
SRT FRET probe	gctactgagatgaaggagacgtgactgtaNH2 Fcttc(Cy3)tctcagtagc	3' Amine	1184
SRT FRET probe	ccaggaagcagttggaggcgtgacgg tNH2 Fcaac(Cy3)gcttcgtgg	3' 2 bases 2''Ome , 3'Amine	1186 1187
h3A4 probe h3A4 invader Capture Sequence	agg agc cac tcc att gga tga agc atg tac aga atc ccc ggt tat tta tgc aga		1188

FIGURE 47P

h3A4 probe h3A4 invader Capture Sequence	gtg gcg tat cac aga caa tga gag cct cct tta tat tcc caa gta taa cac tct aa	1190 1191
Set 2/Set 3 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo h3A4 stacking oligo SRT FRET Oligo	AAC GAG GCG CAC CAC AGA CAA TGA GAG CTCTCATIGICIGGGCG-NH2 cct cct tta tat tcc caa gta taa cac tct aa agctcaatgcatglacagaatccccgg	1192 1193 1195 1196
Set 4 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo SRT FRET Oligo	aac gag gcg cac cac aga caa tga gag ag-NH2 ctc tct cat tgt ctg tgg tgc g-NH2 cct cct tta tat tcc caa gta taa cac tct aa ctc aat gca tgt aca gaa tcc ccg gtt	1197 1198 1200
Set 5 h3A4 probe h3A4 arrestor h3A4 invader SRT FRET probe	aac gag gcg cac cac aga caa tga gag ago t-NH2 agc tct ctc att gtc tgt ggt gcg-NH2 cct cct tta tat tcc caa gta taa cac tct aa FL-caa-c(cy3)g-ctt-cct-ccg	1201 1202 1203 1204
Set 6 h3A4 probe h3A4 arrestor h3A4 invader SRT FRET probe	aac gag gcg cac cac aga caa tga gag agc-NH2 gct ctc tca ttg tct gtg gtg_cgNH2 cct cct tta tat tcc caa gta taa cac tct aa FL-caa-c(cy3)g-ctt-cct-ccg	1205 1206 1207 1208
Set 7/Set 8 h3A4 probe h3A4 probe h3A4 arrestor h3A4 stacking oligo	aac gag gcg cac cac aga caa tga gag a-NH2 aac gag gcg cac cac aga caa tga gag a tct ctc att gtc tgt ggt gcg c-NH2 gct caa tgc atg tac aga atc ccc ggt t	1209 1210 1211 1212

FIGURE 47Q

1213	1214 1215 1216 1217	1218 1219 1220	1222 1223 1224 1225 1225	1228 1229 1230	1232
cct cct tta tat tcc caa gta taa cac tct aa	aac gag gcg cac cac aga caa tga ga-NH2 tct cat tgt ctg tgg tgc gc-NH2 cct cct tta tat tcc caa gta taa cac tct aa gag ctc aat gca tgt aca gaa tcc ccg	AACGAGGCGCACCTCTTATCAGAGCTC AACGAGGCGCACCTCTTATCAGAGCTC-NH2 ttg tgg agg aga tta ttg aga agt gtt gat ta GAGCTCTGATAAGAGGTGCG-NH2	cog tca cgc ctc gcc cca ca - NH2 tgt ggg gcg agg.cg cag cac agg ctg ttg acc atc ata aaa c cuu-uuc-cau-acuuuu-uau-gac-auu-c ctt ttc cag act ttt tat gac att c ctt ttc cag act ttt tat gac	ccg tca cgc ctc gcc cca ca ccg tca cgc ctc gcc cca ca - HEX cag cac agg ctg ttg acc atc ata aaa c cuu-uuc-cau-acu-uuu-uau-gac-auur-c.	ccg tca cgc ctc gcc cca cc - NH2
h3A4 invader SRT FRET Oligo	Set 9 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo SRT FRET Oligo	Set 1/Set 2 h3A4 probe h3A4 probe h3A4 invader h3A4 arrestor SRT	Set 1/ Set 2/ Set 3 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo h3A4 stacking oligo h3A4 stacking oligo SRT FRET	Set 4/Set 5 h3A4 probe h3A4 probe h3A4 invader h3A4 stacking oligo SRT FRET	Set 6/ Set 7/ Set 8 h3A4 probe

FIGURE 47R

h3A4 probe h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo SRT FRET	ccg tca cgc ctc gcc cca cg - NH2 ccg tca cgc ctc gcc cca ct - NH2 tgt ggg gcg agg cg cag cac agg ctg ttg acc atc ata aaa c cag cac agg ctg ttg acc atc ata aaa c cuu-uuc-cau-acu-uuu-uau-gac-auu-c	1233 1234 1235 1236 1237
Set 1 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo SRT	ccg tca cgc ctg atc ata aaa gcc c -NH2 ggg ctt tta tga tca ggc g cag cac agg ctg ttg acc c cac act ttt cca tac ttt tta tg	1238 1239 1240 1241
Set 2 h3A4 probe h3A4 arrestor h3A4 invader h3A4 stacking oligo SRT FRET	aac gag gcg cac cca ttg gat gaa g - NH2 ctt.cat.cca atg ggt gcg c gta cag aat ccc cgg tta ttt atg cag ta ccc.atc.ttc.att.tca.gag	1242 1243 1244 1245
Set 1 h3A5 probe h3A5 invader Capture Sequence	gtg gcg tat cgt gtc taa ttt caa g aat ggg ttt ttc tgg ttg aag aag tcc ttg a	1246 1247
Set 2/Set 3 h3A5 probe h3A5 probe h3A5 arrestor h3A5 invader SRT	AACGAGGCGCACCGTGTCTAATTTCAAG AACGAGGCGCACCGTGTCTAATTTCAAGGG-Pi CTTGAAATTAGACACGGTGCG-NH2 aat ggg tit tic tgg ttg aag aag tcc ttg a	1248 1249 1250
Set 4 h3A5 probe h3A5 arrestor	AACGAGGCGCACCGTGTCTAATTTCAAG CTTGAAATTAGACACGGTGCG-NH2	1252 1253

FIGURE 47S

h3A5 invader h3A5 stacking oligo SRT FRET	aat ggg tit tic tigg tig aag aag toc tig a ggg atc tgt git tot tia caa ggt	1254 1255
Set 5 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	AACGAGGCGCACCGTGTCTAATTTCAAG ctt gaa att aga cac ggt tct c ggt ttt tct ggt tga aga cct tga ggg atc tct gtt tct	1256 1257 1258 1259
Set 6 h3A5 probe h3A5 arrestor h3A5 invader SRT FRET probe	AACGAGGCGCACCGTGTCTAATTTCAAGGG-NH2 CCCTTGAAATTAGACACGGTGCG-NH2 aat ggg ttt ttc tgg ttg aag aag tcc ttg a	1260 1261 1262 1263
Set 7/Set 8 h3A5 probe h3A5 probe h3A5 arrestor h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	aac gag gcg cac cgt gtc taa ttt caa gg-NH2 aac gag gcg cac cgt gtc taa ttt caa gg cct tga aat tag aca cgg tgc gc-NH2 cct tga aat tag aca cgg tgc gc aat ggg ttt tct tgg ttg aag aag tcc ttg a gga tct gtg ttt ctt tac aag gtt tga agg ag	1264 1265 1266 1267 1268
Set 9 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	aac gag gcg cac cgt gtc taa ttt caa-NH2 ttg aaa tta gac acg gtg cgc-NH2 aat ggg ttt ttc tgg ttg aag aag tcc ttg a ggg gat ctg tgt ttc ttt aca agg	1270 1271 1272 1273
Set 10 h3A5 probe	aac gag gcg cac cgt gtc taa ttt ca - NH2	1274

FIGURE 47T

h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	tga aat tag aca cgg tgc gc ggt ttt ct ggt tga aga agt cct tga agg gga tct gtg ttt ct	1275 1276 1277
Set 1 h3A5 probe h3A5 invader Capture Sequence	tgg cgt atc tga ccc ttt ggg aat gaa gag cat aag ttg gaa tca cca cca ta	1278
Set 1 h3A5 probe h3A5 invader Capture Sequence	ata cgg ttg gtc ctc tca agt cta ccc cat tga ttt caa cat ctt tct tgc aac	1280 1281
Set 2/Set 3 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT	aac gag gcg cac gcg tgt cta att tc - NH2 gaa att aga cac gcg tgc gc ggt ttt tct ggt tga aga agt cct tc ccg ggg atc tgt gtt tc	1282 1283 1284 1285
h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	ccg tca cgc ctc gcg tgt cta att tc -NH2 gaa att aga cac gcg agg cg ggt ttt tct ggt tga aga agt cct tc ccg ggg atc tgt gtt tc	1286 1287 1288 1289
Set 1 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT	aac gag gcg cag ttc ata cgt tcc -NH2 gga acg tat gaa ctg cgc cca gca cag gga gtt gac ca cca gca cat tit tcc ata ctt t	1290 1291 1292 1293

Set 2

FIGURE 47U

1294 1295 1296 1297	1298 1299 1300 1301 1303 1303	1305 1306 1307 1308	1309 1310 1312	1313 1315 1315
FIGURE 4/0				
ccg toa cgc ctg ttc ata cgt tcc -NH2 gga acg tat gaa cag gcg cca gca cag gga gtt gac ca cca gca cat ttt tcc ata ctt t	aac gag gcg cac agt tga cct tca aac gag gcg cac agt tga cct tca aac gag gcg cac agt tga cct tca - HEX tga agg tca act gtg cgc gtg atg gcc agc aca ggg c tac gtt ccc cac att ttt c tac gtt ccc cac att ttt c	ccg tca cgc ctc agt tga cct tca tga agg tca act gag gcg gtg atg gcc agc aca ggg c tac gtt ccc cac att ttt c	aac gag gcg cac tcc tct caa gt -NH2 act tga gag gag tgc gc cca ttg att tca aca tct ttc ttg caa ga cta ata gca act ggg aat aat c	cog toa ogo oto too tot caa gt - NH2 act tga gag gag agg cg coa ttg att toa aca tot tto ttg caa ga cta ata goa act ggg aat aat c
h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	Set 1-Set 4 h3A5 probe h3A5 probe h3A5 probe h3A5 arrestor h3A5 arrestor h3A5 invader h3A5 stacking oligo R3A5 STE	Set 5 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	Set 6 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT FRET	Set 7 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT

FIGURE 47V

1317 1318 1319 1320	1321	1323 1324 1325	1326 1327 1328	1330	1332 1333 1334
aac gag gcg cac agt tga cct tc - NH2 tga agg tca act gtg cgc gtg atg gcc agc aca ggg c ata cgt tcc cca cat ttt tc	tgg cgt atc tgg att aaa tct taa aag gac ttt tat tga gag aac gaa tgg atc taa a	AACGAGGCGCACTGGATTAAATCTTAAAAG gac tit lat tga gag aac gaa tgg atc taa a CTTTAAGATTTAATCCAGTGCG-NH2	AACGAGGCGCACTGGATTAAATCTTAAAAG gac ttt tat tga gag aac gaa tgg atc taa a CTTTAAGATTTAATCCAGTGCG-NH2 ctt ctt ggt gtt ttc ca	agg agc cac tca tcc ctt gac t ctt agg gaa atc agg ctc cac tta cgg ta	AACGAGGCGCACCTCATCCCTTGACT AACGAGGCGCACCTCATCCCTTGACT-NH2 AGTCAAGGGAIGAGGGGGG-NH2 ctt agg gaa atc agg ctc cac tta cgg ta
Set 8 h3A5 probe h3A5 arrestor h3A5 invader h3A5 stacking oligo SRT	Set 1 h3A7 Probe h3A7 Invader Capture Oliog	Set 2 h3A7 Primary Probe h3A7 Invader h3A7 Arrestor SRT FRET	Set 3 h3A7 Primary Probe h3A7 Invader h3A7 Arrestor h3A7 Stacking Oligo SRT FRET	Set 4 h3A7 Probe h3A7 Invader oligo Capture Oligo	Set 5/Set 6 h3A7 Primary Probe h3A7 Primary Probe h3A7 Arrestor h3A7 Invader oligo

FIGURE 47W

1336 1338 1339 1340 1341	1343 1344 1345 1346 1347	1348	1350 1351 1352	1353 1354 1355 1356
aac gag gcg cac cto atc cct tga c-NH2 gtc_aag gga tga gcg_c-NH2 ctt agg gaa atc agg ctc cac tta cgg ta tca gcc ttt aga aca atg ggt ttt tct gtt ag3' tca gcc ttt aga aca atg ggt ttt tct g ctc agc_ctt tag aac aat ggg ttt tct ct ctc agc_ctt tag aac aat ggg ttt tct ct ctc agc_ctt tag aac aat ggg ttt ttc t	aac gag gcg cac ctc atc cct tga-NH2 aac gag gcg cac ctc atc cct tga c tca agg gat gag gtg cgc-NH2 ctt agg gaa atc agg ctc cac tta cgg ta ctc agc ctt tag aac aat ggg ttt ttc tgt tag	ata cgg ttg gta aag taa ttt gag gt gaa gcc cgt ctt cat ttc agg gtt cta ttt c	AACGAGGCGCACGTAAAGTAATTTGAGGT gaa gcc cgt ctt cat ttc agg gtt cta ttt c ACCTCAAATTACTTTACGTGCG-NH2	AACGAGGCGCACGTAAAGTAATTTGAGGT gaa gcc cgt ctt cat ttc agg gtt cta ttt c ACCICAAATTACTTTACGTGCG-NH2 ctc tgg tgt tct ggg
Set 7 - Set 10 h3A7 Primary Probe h3A7 Arrestor h3A7 Invader oligo h3A7 Stacking Oligo h3A7 Stacking Oligo h3A7 Stacking Oligo h3A7 Stacking Oligo SRT FRET	Set 11 h3A7 Primary Probe h3A7 Primary Probe h3A7 Arrestor h3A7 Invader oligo h3A7 Stacking Oligo SRT FRET	Set 1 h3A7 Probe h3A7 Invader Capture Sequence	Set 2 h3A7 Primary Probe h3A7 Invader h3A7 Arrestor SRT FRET	Set 3 h3A7 Primary Probe h3A7 Invader h3A7 Arrestor h3A7 Stacking Oligo

FIGURE 47X

	FIGURE 4/X	
set 1 h3A7 probe h3A7 arrestor h3A7 stacking oligo SRT	cog toa cgo oto gto ata aat aco co - NH2 ggg gto tit atg acg agg cg goc ago ata ggo tot tga cao aga ctt tit tat act tit tat aac att c	1357 1358 1359 1360
Set 2 - Set 4 h3A7 probe h3A7 probe h3A7 probe h3A7 arrestor h3A7 arrestor h3A7 stacking oligo SRT	aac gag gcg cac gtc ata aat acc cc -NH2 aac gag gcg cac gtc ata aat acc cc aac gag gcg cac gtc ata aat acc cc - HEX ggg gta ttt atg acg tgc gc gcc agc ata ggc tgt tga cac aga ctt ttc tat act ttt tat aac att c	1361 1362 1363 1364 1365
Set 1 h3A7 probe h3A7 arrestor h3A7 invader h3A7 stacking oligo SRT	cog tca cgc ctc gat taa atc tta aaa gct t - NH2 aag ctt tta aga ttt aat cga ggc g gac ttt tat tga gag aac gaa tgg atc taa tgc ctt ggt gtt ttc cac aaa g	1367 1368 1369
Set 2 h3A7 probe h3A7 arrestor h3A7 invader h3A7 stacking oligo SRT	aac gag gcg cac gat taa atc tta aaa gct t -NH2 aag ctt tta aga ttt aat cgt gcg c gac ttt tat tga gag aac gaa tgg atc taa tgc cft ggt gtt ttc cac aaa g	1371 1372 . 1373 1374
Set 1 h3A7 probe h3A7 arrestor	ccg tca cgc ctg tca tcc ctt g - NH2	1375

FIGURE 47Y

FIGURE 4 / Y 1377 1378	1379 1380 1381 1382	- NH2 1383 1384 1385 1386 1386 1386 1386 1386 1386 1386 1386	1387 a	CC 1389 1390 1390 1391 1391 1391 1392 1392	1393
gga aat cag gct cca ctt acg gtc a a ct cag cct tta gaa caa tg	ceg tca egc ctc taa agt aat ttg agg tc -NH2 gac ctc aaa tta ctt tag agg cg egt ctt cat ttc agg gtt cta ttt ga tct ggt gtt ctg gg	aac gag gcg cac taa agt aat ttg agg tc - NH2 gac ctc aaa gga ctt tag tgc gc cgt ctt cat ttc agg gtt cta ttt ga tct ggt gtt ctg gg	tgg-cgt-atc-tag-gct-ttg-ctt-cc ttc atg tag tca ggg tca tag aca att aag a	AACGAGGCGCACTAGGCTTTGCTTCC GGAAGCAAAGCCTAGTGCG-NH2 gga agc aaa gcc tag tgc gc-NH2 ttc atg tag tca ggg tca tag aca att aag a	aac gag gcg cac tag gct ttg ctt ccc-NH2 ggg aag caa agc cta gtg cgc-NH2
h3A7 invader h3A7 stacking oligo SRT FRET	Set 1 h3A7 probe h3A7 arrestor h3A7 invader h3A7 stacking oligo SRT	Set 2 h3A7 probe h3A7 arrestor h3A7 invader h3A7 stacking oligo SRT	Set 1 r4A1 Probe r4A1 invader Capture Sequence	Set 2 r4A1 Primary Probe r4A1 Arrestor r4A1 Arrestor r4A1 Invader FRET Probe 1	Set 3 r4A1 Primary Probe

FIGURE 47Z

1396 1397 1398 1399	1400 1401 1402 1403	1405 1406 1407 1408 1410 1411	1413 1414	1415 1416 1417
aac gag gcg cac tag gct ttg ctt c-NH2 gaa gca aag cct agt gcg c ccc aga acc atc gag gaa agg c ttc atg tag tca ggg tca tag aca att aag a	aac gag gcg cac tag gct ttg ctt-NH2 aag caa agc cta gtg cgc-NH2 ttc atg tag tca ggg tca tag aca att aag a ccc cag aac cat cga gga aag g ccc cag aac cat cga gga aag g	aac gag gog cac tag got ttg ct-NH2 aac gag gog cac tag got ttg ct - HEX aac gag gog cac tag got ttg ct agc aaa goc tag tgc gc-NH2 agc aaa goc tag tgc gc ttc atg tag tgc gc ttc atg tag tag aca att aag a ttc cca gaa cca tcg agg aaa gg	ata cgg ttg gtc ttg acc tgc c agg aga tat gtt gaa aga ttt cta tag agg ac	AACGAGGCGCACGTCTTGACCTGCC GGCAGGTCAAGACGTGCG-NH2 agg aga tat gtt gaa aga ttt cta tag agg ac
Set 4 r4A1 Primary Probe r4A1 Arrestor r4A1 Stacker r4A1 Invader SRT FRET Probe 1	Set 5 r4A1 Primary Probe r4A1 Arrestor r4A1 Invader r4A1 Stacker r4A1 Stacker r4A1 Stacker FRET Probe 1	Set 6 r4A1 Primary Probe r4A1 Primary Probe r4A1 Arrestor r4A1 Arrestor r4A1 Invader r4A1 Stacker r4A1 Stacker FRET Probe 1	Set 1 r4A1 Probe r4A1 Invader Capture Sequence	Set 2 r4A1 Primary Probe r4A1 Arrestor r4A1 Invader

FIGURE 47AA

SRT FRET Probe 1

Set 3 r4A1 Primary Probe r4A1 Arrestor r4A1 Invader SRT FRET Probe 1	AACGAGGCGCACGTCTTGACCTGC-Pi GGCAGGTCAAGACGTGCG-NH2 agg aga tat gtt gaa aga ttt cta tag agg ac	1418 1419 1420
Set 1 r4A1 Probe r4A1 Invader	tgg cgt atc tta gat gga gta agg a att cct cat aat tca aaa ggg act tag tag gt	1421 1422
Set 2 r4A1 Primary Probe r4A1 Arrestor SRT FRET Probe 1	AACGAGGCGCACTTAGATGGAGTAAGGA ICCITACICCAICIAAGIGCG-NH2	1423 1424
Set 1 r4A1 Primary Probe r4A1 Arrestor r4A1 Invader SRT FRET Probe 1	aac gag gcg cac tgg ata ccc ttg gg-NH2 ccc aag ggt atc cag tgc gc-NH2 ggt gga gac cat aaa tgg aga gtg tga cta	1425 1426 1427
Set 1 r4A2 Probe r4A2 Arrestor r4A2 Invader SRT FRET Probe 1	aac gag gcg cac agg tgt ctg gag taa aag-NH2 ctt tta ctc cag aca cct gtg cgc -NH2 gtc cac gca caa gct ggg ac	1428 1429 1430
Set 1 r4A2 Probe r4A2 Arrestor r4A2 Invader r4A2 stacking oligo SRT	aac gag gcg cac aga agg ccc ctt-NH2 aag ggg cct tct gtg cgc-NH2 cct tga aca gca cca gaa ata gac tga gca c gga aga acc cag aga cac cat cc	1431 1432 1433 1434

FIGURE 47AB

FRET Probe 1

Set 2 r4A2 Probe r4A2 Arrestor r4A2 Invader SRT FRET Probe 1	oog toa ogo oto aga agg ooc ott-NH2 aag gg<u>g oot</u> tot gag gog -NH2 oot tga aca goa ooa gaa ata gao tga goa o	1435 1436 1437
Set 3 r4A2 Probe r4A2 Arrestor r4A2 Invader SRT FRET Probe 1	aac gag gog cac aga agg ccc ctt g-NH2 caa ggg go <u>c ttc tgt gog c</u> -NH2 cct tga aca gca cca gaa ata gac tga gca c	1438 1439 1440
Set 4 r4A2 Probe r4A2 Probe r4A2 Probe r4A2 Arrestor r 4A2 Arrestor r4A2 Invader SRT	aac gag gcg cac aga agg ccc ctt gg-NH2 aac gag gcg cac aga agg ccc ctt aac gag gcg cac aga agg ccc ctt - HEX cca agg ggc ctt ctg tgc gc-NH2 aag ggg cct tct gtg cgc cct tga aca gca cca gaa ata gac tga gca c	1441 1442 1443 1444 1446
Set 1 r4A3 Probe r4A3 Arrestor r4A3 Invader SRT FRET Probe 1	aac gag gcg cac ttg aca gag tcc gc-NH2 gcg gac tct gtc aag tgc gc-NH2 gct tct ccc att tgt cta gca tta taa	1447 1448 1449
Set 2 r4A3 Probe r4A3 Arrestor r4A3 Invader r4A3 stacking oligo SRT FRET Probe 1	aac gag gcg cac ttg aca gag tcc g-NH2 cag act ctg tca agt gcg c-NH2 gct tct ccc att tgt cta gca tta taa cca tga ttt tga cat agg gtt tga gga tg	1450 1451 1452 1453

FIGURE 47AC

	1454 1455 1456 1457 1459 1460	1461	1463 1464 1465	1466	1468 1469 1470 1471	1473
FIGURE 47AC						,
	aac gag gcg cac ttg aca gag tcc-NH2 aac gag gcg cac ttg aca gag tcc aac gag gcg cac ttg aca gag tcc - HEX gga ctc tgt caa gtg cgc-NH2 gga ctc tgt caa gtg cgc gct tct ccc att tgt cta gca tta taa gct tct ccc att tgt cta gca tta taa gcc atg att ttg aca tag ggt ttg agg atg	cgg agc ctc tgc ggt cat caa g tgg ata act gca tca gtg tat ggc att tta a	gtg-gcg-tat-ctg-cgg-tca-tca-ag gtg-gcg-tat-ctg-cgg-tca-tca-a tgg ata act gca tca gtg tat ggc att tta a	tg-gog-tat-ctg-ogg-tca-tca-a tgg ata act goa tca gtg tat ggc att tta a	aac-gag-gcg-cac-ctg-cgg-tca-tca-a ttg-atg-acc-gca-ggt-gcg-cc-NH2 ttg-atg-acc-gca-ggt-gcg-cc-Pi ttg-atg-acc-gca-ggt-gcg-cc-OH ttg-atg-acc-gca-ggt-gcg-cc-OH	aac-gag-gcg-cac-ctg-cgg-tca-tca-a
	Set 3 r4A3 Probe r4A3 Probe rCYP 4A3 Probe r4A3 Arrestor rCYP 4A3 Arrestor r4A3 Invader r4A3 stacking oligo SRT FRET Probe 1	Set 1 r281 probe r281 invader Capture Sequence	Set 2/ Set 3 r2B1 probe r2B1 probe r2B1 invader Capture Sequence	Set 4 r2B1 probe r2B1 invader Capture Sequence	Set 5 - Set 7 r2B1 probe r2B1 arrestor r2B1 arrestor r2B1 invader SRT	Set 8 r2B1 probe

FIGURE 47AD

1474 1475 1476	1477 1478 1479	1480 1481 1482	1483 1484 1485	1486 1487 1488	1489 1490 1491
ttg-atg-acc-gca-ggt-gcg-cc- Pi	aac-gag-gcg-cac-ctg-cgg-tca-tca-a-NH2	ggo-aac-gag-gca-cac-ctg-cgg-tca-tca-ag-Pi	aac gag ggg cac ctg cgg tca tca ag-NH2	aac gag gcg cac ctg cgg tca tca agg-NH2	atg acg tga cag acc tgc ggt cat caa g-NH2
tgg ata act gca tca gtg tat ggc att tta a	ttg-atg-acc-gca-ggt-gcg-NH2	ttg-atg-acc-gca-ggt-gcg-cc- Pi	ctt gat gac cgc agg tgc c-NH2	cct tga tga cg cag gtg cg -NH2	cft gat gac cgc agg tct gt -NH2
ggg ttg gta gcc tgt gtg agc cga t	tgg ata act gca tca gtg tat ggc att tta a	tgg ata act gca tca gtg tat ggc att tta a	tgg ata act gca tca gtg tat ggc att tta a	tgg ata act gca tca gtg tat ggc att tta a	tgg ata act gca tca gtg tat ggc att tta a
ttg-atg-acc-gca-ggt-gcg-cc- Pi	aac-gag-gcg-cac-ctg-cgg-tca-tc	ggc-aac-gag-gca-cac-ctg-cgg-tc	aac gag ggg cac	aac gag gcg cac ctg cgg tca tca a	atg acg tga cag acc tgc ggt cat c
tgg ata act gca tca gtg tat ggc a	ttg-atg-acc-gca-ggt-gcg-NH2	ttg-atg-acc-gca-ggt-gcg-cc -Pi	ctt gat gac cgc ag	cct tga tga ccg cag gtg cg -NH2	ctt gat gac cgc agg tct gt -NH2
ggg ttg gta gcc tgt gtg agc cga t	tgg ata act gca tca gtg tat ggc s	tgg ata act gca tca gtg tat ggc a	tgg ata act gca tc	tgg ata act gca tca gtg tat ggc att	tgg ata act gca tca gtg tat ggc at
r2B1 arrestor r2B1 invader r2B1 stacker SRT FRET	Set 9 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET	Set 10 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET	Set 11 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET	Set 12 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET	Set 13 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET

FIGURE 47AE

1492 1493 1494	1495 1496 1497	1498 1499 1500	1501 1502 1503 1504	1505 1506 1507 1508	
aac gag gcg cac ctg agg tca tca a-NH2 ttg atg acc tca ggt gcg -NH2 tgg ata act gca tca gtg tat ggc att tta a	cag toa cgt ctc ctg cgg toa toa ag-NH2 ctt gat gac cgc agg aga cg-NH2 tgg ata act goa toa gtg tat ggc att tta a	cag toa ogt oto act gog gto ato aag-NH2 gtg gat aac tgo ato agt gta tgg cat ttt o ott gat gac ogo agt gag acg-NH2	cag toa ogt oto act gog gto ato aa-NH2 ttg atg acc goa gtg aga cg-NH2 gtg gat aac tgo ato agt gta tgg cat ttt o ggg ttg gta gcc tgt gtg ago oga t	cag toa cgt ctc act gcg gtc atc a-NH2 tga tga ccg cag tga gac g-NH2 gfg gat aac tgc atc agt gta tgg cat ttt c agg gtt ggt agc ctg tgt gag ccg a	can tea out ofe act one ate ate.
Set 14 12B1 probe 12B1 arrestor 12B1 invader SRT FRET	Set 15 r2B1 probe r2B1 arrestor r2B1 invader SRT FRET	Set 16 r2B1 probe r2B1 invader r2B1 arrestor SRT FRET	Set 17 r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker SRT	Set 18 r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker SRT FRET	Set 19

FIGURE 47AF

1510 1511 1512	1513 1514 1515 1516	1517 1518 1519 1520	1521 1522 1523 1524	1525 1526 1527 1528 1529	1530 1531
					ı
z #c	o #	tH2	# 12	# c	,a aag
ctt gat gac cgc agt gag acg- NH2 gtg gat aac tgc atc agt gta tgg cat ttt c ggt tgg tag cct gtg tga gcc gat c	cag tca cgt ctc act gcg gtc at-NH2 atg acc gca gtg aga cg-NH2 gtg gat aac tgc atc agt gta tgg cat ttt c caa ggg ttg gta gcc tgt gtg agc c	cog tca cgc cto act gog gtc atc a-NH2 tga tga ccg cag tga ggc g-NH2 gtg gat aac tgc atc agt gta tgg cat ttt c agg gtt ggt agc ctg tgt gag ccg a	cog tca cgc ctc act gcg gtc atc-NH2 gat gac cgc agt gag gcg-NH2 gtg gat aac tgc atc agt gta tgg cat tit c aag ggt tgg tag ceg gtg tg	cog tca cgc ctc act gog gtc at-NH2 cog tca cgc ctc act gog gtc at atg acc gca gtg agg cg-NH2 gtg gat aac tgc atc agt gta tgg cat ttt c caa ggg ttg gta gcc tgt gtg agc c	atg gtg tct ttg gtg act ctg tgt ggt aca aac-gag-gac-cac-tcc-aat-agg-gac-aag
gc agt ga gc atc agt g xt gtg tga g	c act gog ; gig aga c; gc atc agt ; ta gcc tgt ;	tc act gog zag tga go yc atc agt g yc ctg tgt g	tc act gcg agt gag g yc atc agt g	ccg tca cgc ctc act gcg gtc al· ccg tca cgc ctc act gcg gtc at atg acc gca gtg agg cg- NH2 gtg gat aac tgc atc agt gta tgg caa ggg ttg gta gcc tgt gtg ag	gtg act ctg
gat gac c gat aac te tgg tag cc	g tca cgt cl Lacc gca gat aac tc ggg ttg g) tca cgc c L íga ccg c gat aac tc ggt ggt ag	t ca cgc c Lgac cgc gat aac tg ggt tgg ta) tca cgc c) tca cgc c Lacc gca gat aac tc ggg ttg g	gtg tct ttg
ct 91g	gag grg gag	ccc grg grg agg	ccc grg	cocc	atg
r2B1 arrestor r2B1 invader r2B1 stacker SRT FRET	Set 20 r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker SRT	Set 21 r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker SRT	Set 22 r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker	Set 23 r2B1 probe r2B1 probe r2B1 arrestor r2B1 invader r2B1 stacker SRT	Set 1 r2B1 invader r2B1 probe
7281 ii 7281 ii 7281 s SRT FRET	Set 20 r2B1 probe r2B1 arrestt r2B1 invade r2B1 stacke SRT	Set 21 r2B1 probe r2B1 arrestr r2B1 invade r2B1 stacke SRT	Set 22 r2B1 probe r2B1 arrestr r2B1 invade r2B1 stacke	Set 23 r2B1 probe r2B1 probe r2B1 arrestt r2B1 invade r2B1 stacke SRT FRET	Set 1 r2B1 invade r2B1 probe

FIGURE 47AG

r2B1 arrestor SRT FRET	ctt-gtc-cct-att-gga-gtg-cgc-c	1532
Set 1 r2B1 probe r2B1 invader Capture Sequence	gcg gcg tac agc cgg tgt gag c cat ttt act gcg gtc atc aag ggt tgg tc	1533 1534
r2B1 probe r2B1 invader Capture Sequence	tgg cgt atg agc cgg tgt gag c cat ttt act gcg gtc atc aag ggt tgg tc	1535 1536
Set 1 r2B2 invader r2B2 probe r2B2 arrestor SRT FRET	gga tga ctg cat cag tgt atg gca ttt tgc aac-gag-gcg-cac-gta-cga-tca-tca-agg cct-tga-tga-tcg-tac-gtg-cgc-c-NH2	1537 1538 1539
Set 1 r2B2 invader r2B2 probe r2B2 stacker r2B2 invader stacker	atg gtg tot ttg gtg act ctg tgt ggt aac tgg cgt atg acc aat tgg ggc aa gat ctg caa atc tct gaa tct cgt gga tg tct tgg aga gca ggt acc ctc gga ac	1540 1541 1542 1543
Set 2 r2B2 probe r2B2 invader r2B2 stacker r2B2 invader stacker	tgg cgt atg acc aat tgg ggc aag atg gtg tct ttg gtg act ctg tgt ggt aac atc tgc aaa tct ctg aat ctc gtg gat ga tct tgg aga gca ggt acc ctc gga ac	1544 1545 1546 1547
Set 3 r2B2 probe r2B2 probe r2B2 arrestor r2B2 invader SRT FRET	aac-gag-gcg-cac-acc-aat-tgg-ggc-aag aac gac gcg cac acc aat tgg ggc aag ctt-gcc-cca-att-ggt-ctgc-cc- NH2 atg gtg tct ttg gtg act ctg tgt ggt aac	1548 1549 1550 1551

FIGURE 47AH

1552 1553 1554	1555 1556 1557 1558	1559 1560 1561	1562 1563 1564	1565 1566 1567 1568	1569
	•				
aac-gag-gcg-cac-acc-aat-tgg-ggc-aag-Pi ctt-gcc-cca-att-ggt-gtg-cgc-c-Pi atg gtg tct ttg gtg act ctg tgt ggt aac	ctt gcc cca att ggt gtg cg-NH2 aac-gag-gcg-cac-acc-aat-tgg-ggc-aag-NH2 atg gtg tct ttg gtg act ctg tgt ggt aac atc tgc aaa tct ctg aat ctc gtg gat ga	ggc-aac-gag-gca-cac-caa-ttg-ggg-caa-g ctt-gcc-cca-att-ggt-gtg-cgc-c-NH2 atg gtg tct ttg gtg act ctg tgt ggt aac	aac gag gog cac acc aat tgg ggc aag atc-NH2 gat ctt gcc cca att ggt gtg cg-NH2 atg gtg tct ttg gtg act ctg tgt ggt aac	aac gag gog cac acc aat tog ggc aag-NH2 ctt gcc cga att ggt gtg cg-NH2 atg gtg tot ttg gtg act ctg tgt ggt aac atc tgc aaa tct ctg aat ctc gtg gat ga	cag tca cgt ctc atg gtg gcc tgt g-NH2
Set 4 (282 probe (282 arrestor (282 invader SRT FRET	Set 5 r2B2 arrestor r2B2 probe r2B2 invader r2B2 stacker SRT FRET	Set 6 r2B2 probe r2B2 arrestor r2B2 invader SRT FRET	Set 7 r2B2 probe r2B2 arrestor r2B2 invader SRT FRET	Set 8 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker SRT	Set 9 r2B2 probe

FIGURE 47AI

r2B2 invader r2B2 arrestor SRT FRET	gta tgg cat ttt ggt acg atc atc aag ggc cac agg cca cca tga gac g-NH2	1570 1571
Set 10 r2B2 probe r2B2 invader r2B2 arrestor r2B2 stacker SRT FRET	cag tca cgt ctc aga gcc aat cac ctg-NH2 cga tca tca agg gat ggt ggc ctg tgc cag gtg att ggc tct gag acg -NH2 atc aat ctc ctt ttg gac ttt ctc tgc g	1572 1573 1574 1575
Set 11 r2B2 probe r2B2 invader r2B2 arrestor r2B2 stacker SRT FRET	cag toa cgt oto aga goo aat cac ot-NH2 cga toa toa agg gat ggt ggo otg tgo agg toa ttg got otg aga cg-NH2 gat caa tot cot ttt gga ott tot otg o	1576 1577 1578 1579
Set 12 r2B2 probe	FAM-cag toa ogt otc aga goc aat cac ct-NH2	1580
Set 13 / Set 14 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker r2B2 stacker SRT FRET	cag toa ogt oto aga goo aat cac c-NH2 ggt gat tgg oto tga gac g-NH2 oga toa toa agg gat ggt ggc otg tgc gat caa tot cot tit gga cit tot oc tga toa atc toc tit tgg act tot oc	1581 1582 1583 1584 1585
Set 15 r2B2 probe r2B2 arrestor r2B2 stacker r2B2 invader SRT FRET	cag tca cgt ctc aga gcc aat cac-NH2 gtg att ggc ict gag acg -NH2 ctg atc aat ctc ctt ttg gac ttt ctc tgc g cga tca tca agg gat ggt ggc ctg tgc	1586 1587 1588 1589

FIGURE 47AJ

1590 1591 1592 1593	1594 1595 1596 1597	1598 1599 1600	- 1602 1603 1604 1605	1606 1607 1608 1609 1610
ct-NH2	atg-NH2 12 9c	ct-NH2 2 gc	o-NH2 go c	NH2 gc igc g
cag tca cgt ctc aga ggc aat cac ct-NH2 agg tga ttg cct ctg aga cg-NH2 cga tca tca agg gat ggt ggc ctg tgc gat caa tct cct ttt gga ctt tct ctg c	cag toa ogt oto aga ggo aat oac otg-NH2 cag gtg att goc tot gag acg-NH2 oga toa toa agg gat ggt ggo otg tgo atc aat oto ott ttg gao ttt oto tgo g	cog tca cgc ctc aga gcc aat cac ct-NH ² agg tga ttg gct ctg agg cg-NH ² cga tca tca agg gat ggt ggc ctg tgc gat caa tct cct ttt gga ctt tct ctg c	ccg tca cgc ctc aga gcc aat cac c-NH2 ggt gat tgg ctc tga ggc g-NH2 cga tca tca agg gat ggt ggc ctg tgc tga tca atc tcc ttt tgg act ttc tct gc	ccg tca cgc ctc aga gcc aat cac-NH2 ccg tca cgc ctc aga gcc aat cac gtg att ggc tct gag gcg-NH2 cga tca tca agg gat ggt ggc ctg tgc ctg atc aat cat ct ttg gac ttt ctc tgc g
Set 16 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker SRT FRET	Set 17 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker SRT FRET	Set 18 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker SRT FRET	Set 19 r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker SRT FRET	Set 20-21 r2B2 probe r2B2 probe r2B2 arrestor r2B2 invader r2B2 stacker

FIGURE 47AK

Set 22	MA/+ ANDDIA	
r282 probe r282 invader r282 arrestor SRT FRET	cag tca cgt ctc atg gtc aaa gta ctg tgg-NH2 gga agt gct cag gat tga agg tgt ctg gc cca cag tac.ttt gac.cat.gag.acg-NH2	1611 1612 1613
Set 23 r2B2 probe r2B2 arrestor r2B2 invader SRT FRET	aac gag gcg cac atg gtc aaa gta ctg tgg-NH2 cca cag tac ttt gac cat gtg cgc -NH2 gga agt gct cag gat tga agg tgt ctg gc	1614 1615 1616
r2B2 probe r2B2 invader	cat acg gtt ggg cct gtg aga gc cat ttt ggt acg atc atc aag gga tgg tc	1617 1618
		•
r3A1 probe	agg agc cac ggg tcc caa atc El sam ant cat matter caa atc	1619
r3A1 invader	i E-ayy ayo cac yyy ico caa aic toc oot git tot tga aaa gto cat gtg tga	1621
r3A1 probe	F-tog ogt agt ogg gtc cca aat c	1622
r3A1 probe	cat-ctt-cgc-gga-cgg-gtc-cca-aat-c	1623
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gcg-cc-NH2	1624
r3A1 probe	aac-gag-gcg-cac-cgg-gtc-cca-aat-c-NH2	1625
r3A1 probe	cat-ctt-cgc-gga-cgg-gtc-cca-aat-c - NH2	1626
r3A1 arrestor	gga ttt ggg acc cgt ccg cga - NH2	1627
r3A1 arrestor	ggattirgggtatcc-cgr-ccg-cg -wnz oos ttt ood see eet eet ee - NH2	1528
r3A1 arrestor	aga tit ada acc cat cca - NH2	1630
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gcg-c-NH2	1631
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gcg-NH2	1632
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gc-NH2	1633
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gcg-cct-NH2	1634
r3A1 arrestor	gat-ttg-gga-ccc-ggt-gcg-cct-c-NH2	1635
r3A1 probe		
r3A1 probe	aac gag gcg cac cgg gtc cca aat c-Pi	1636

FIGURE 47AL

r3A1 invader	tcc cct att tct taa aaa atc cat ata taa	1637
-2 A 4 5-54 C		000
	ade gag geg care egg gic cervana	1038
r3A1 arrestor	gat ttg gga ccc ggt gcg-NH2	1639
r3A1 probe	aac gag gcg cac cgg gtc cca aat c-NH2	1640
r3A1 arrestor	ggaitt ggg acc cgg tgc gc-NH2	1641
r3A1 probe	aac gag gcg cac cgg gtc oca aat-NH2	1642
r3A1 arrestor	att tgg gac ccg gtg cgc-NH2	1643
r3A1 stacker	ccg tag agg agc acc agg acg	1644
r3A1 probe	aac gag gcg cac cgg gtc cca aa-NH2	1645
r3A1 arrestor	ttt ggg acc cgg tgc gc-NH2	1646
r3A1 stacker	toc gta gag gag cac cag ga	1647
r3A1 probe	cag tca cgt ctc cgg gtc cca aa-NH2	1648
r3A1 arrestor	ttt ggg acc cgg aga cg-NH2	1649
r3A1 stacker	tcc gta gag gag cac cag ga	1650
r3A1 probe	ccg tca cgc ctc cgg gtc cca aa-NH2	1651
r3A1 arrestor	ttt ggg acc cgg agg cg-NH2	1652
r3A1 stacker	t<u>cc. g</u>ta gag gag cac cag ga	1653
r3A1 stacker	toe gta gag gag cae cag ga	1654
r3A1 probe	aac gag gcg cac cgg gtc cca-NH2	1655
r3A1 arrestor	tgg gac ccg gtg cgc-NH2	1656
r3A1 probe	ccg tca cgc ctc cgg gtc cca-NH2	1657
r3A1 arrestor	tgg gac ccg gag gcg-NH2	1658
r3A1 stacker	aat ccg tag agc acc agg	1659
r3A1 probe	aac gag gcg cac cgg gtc cca	1660
r3A2 invader	tto cit git tot taa aaa tto cat gto taa	1661
r3A2 invader	att ttt oga tac ttt tta tag cac toc atc	1662
r3A2 probe	tgg cgt atc tgg gtt cca agt c	1663
r3A2 probe	aac gag gcg cac gtc aaa tct ccc taa	1664
r3A2 probe	aac-gag-gcg-cac-tgg-gtt-cca-agt-c	1665
r3A2 arrestor	tta ggg aga ttt gac gtg cgc c - NH2	1666
r3A2 arrestor	gac-ttg-gaa-ccc-agt-gcg-cc-NH2	1667
r3A2 probe	aac gac gcg cac tgg gtt cca agt c	1668
r3A2 probe	aac-gag-gcg-cac-tgg-gtt-cca-agt-c-Pi	1669
r3A2 arrestor	ga <u>c ttg gaa ccc agt gcg</u> -NH2	1670
r3A2 probe	aac gag gcg cac tgg gtt cca agt cg-NH2	1671
r3A2 arrestor	cga ctt gga acc cag tgc gc-NH2	1672
r3A2 probe	aac gag gcg cac aac cat caa gtt cta ta-NH2	1673

FIGURE 47AM

gga atc gtc act act gac cct ttg ggt ata aac ac 1674 tct ttt tta cag act ctc tca agt cta tta cc 1675 testor tat aga act toa tog ttg toc ac-NH2	aac gag gcg cac aac cat caa gtt cta-NH2	acker tat cit itt tac aga ctc tct caa gtc tat tac c 1678 restor tag aac ttg atg ott ctc cae gtc -NH2	cag tca cgt ctc ctc ggc agg gc-NH2	vader cac aat atc gta ggt agg agg tgc ctt aa 1681	33B	obe cag toa cgt ctc ctc ggc agg g-NH2		restor ccc tgc cga gga gac g-NH2 1685	obe cag tca cgt ctc ctc ggc agg-NH2 1686	acker gcc cca tog atc tcc tcc 1687	CCT	cag tca cgt ctc ctc ggc ag-NH2	ggc ccc atc gat ctc ctc	restor ctg ccg agg aga aga cg-NH2 1691	obe ccg tca cgc ctc ctc ggc agg-NH2 1692	restor cct gcc gag gag gcg-NH2 1693	acker gcc cca tcg atc tcc tcc 1694	obe ccg tca cgc ctc ctc ggc agg	1 probe ccg tca cgc ctc ggc ttg tgt gtt c-NH2	cca aga tag att cag aga agc atc	and the ato and ale cet	gaa cac aca ago cga ggc g	1-1 probe ccg tca cgc ctc gcc ttt gtt tgg-NH2	cca aac aaa ggc gag gcg	ggg caa cat tga cat aaa gtg ttt gog tac tct c	甘	ccg tca cgc ctc gcc ttt gtt tg-NH2	caa aca aag gog agg cg	stacker ggt tcg aat tcc atg tca tc	CHIM of one one for for one one	on probe and gag gdg dad gdt cct gga aga tg-NH2 OH arrestor cat ctt cca gga gcg tgc gcc -NH2 1708
r3A2 invader r3A2 stacker r3A2 arrestor	r3A2 probe	r3A2 stacker r3A2 arrestor	r3A2 probe	r3A2 invader	r3A2 arrestor	r3A2 probe	r3A2 stacker	r3A2 arrestor	r3A2 probe	r3A2 stacker	r3A2 arrestor	r3A2 probe	r3A2 stacker	r3A2 arrestor	r3A2 probe	r3A2 arrestor	r3A2 stacker	r3A2 probe	hICAM-1 probe	hICAM-1 invader	hICAM-1 stacker	hICAM-1 arrestor	hVCAM-1 probe	hVCAM-1 arrestor	hVCAM-1 invader	hVCAM-1 stacker	hVCAM-1 probe	hVCAM-1 arrestor	hVCAM-1 stacker	Odora HOOVOA	hGAPDH arrestor

FIGURE 47AN

hGAPDH invader	cac ttg att ttg gag gga tct ca	1709
Secondary system oligos	ligos	
Capture Oligo Capture Oligo Capture Oligo Capture Oligo Capture Oligo	aaa agt ggc tcc t-(biotin)c aaa aga ggc tcc gct-(biotin) c aaa aga tac gcc gct-(biotin) c aaa aga tac gcc aca gct-(biotin) c aaa acc aac cgt atg aac t-(biotin) c aaa atc ata cgc cac t-(biotin)c	1710 1711 1712 1713 1714
SRT SRT SRT SRT SRT SRT SRT SRT SRT SRT	cgg-agg-aag-cag-ttg-gtg-tgc-ctc-gtt-gcc-tt-NH2 cgg aag aag cag ttg gtg ccc ctc gtt aa-NH2 cgg aag aag cag ttg gtg ccc ctc gtt aa-NH2 cgg aag aag cag ttg gtg cgc ctc gtt aa-NH2 cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gtg cgc ctc gtt aa cgg aag aag cag ttg gag gcg tga cgg t-NH2 cgg aag aag cag ttg gag gcg tga cgg t-NH2 cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t cgg aag aag cag ttg gag gcg tga cgg t	1716 1717 1718 1719 1720 1721 1723 1724 1725 1726 1727
FRET probe FRET probe FRET probe FRET probe FRET probe	FL-caa c(cy3)gc ttc ctc FL-caa c(cy3)gc ttc ctc FL-caa-c(cy3)g-ctt-cct-ccg FL-caa-c(cy3)g-ctt-cct-ccg- <u>uu</u> FL-caa-c(cy3)g-ctt-cct-ccg- <u>uuu-u</u> FL-caa-c(cy3)g-ctt-cct-ccg-uuu-u	1730 1731 1732 1733 1734

FIGURE 47AO

Oligo sequence descriptions: 5' to 3' direction, 2'-Ome nts are bolded and underlined, internal modifications are defined in (), ASR of primary probes are underlined C18ddC = C18 linker+dideoxy C, ddC = dideoxy C, FI = Fluorescein

Oligo Type	Oligo Seguesce	ON CI CHO
HIMAN II -2		SEG ID NO
Human IL-2 Probe	FI- CGAAATTAATACGCCTTCTTGGGCATGTAC -C18ddC	1736
Human IL-2 Probe	CGAAATTAATACGCCTTCTTGGGCATGTAC -C18ddC	1737
Human IL-2 Invader	CTGAAGATGTTTCAGTTCTGTG- ddC	1738
Human IL-2 Invader	GAAGATGTTTCAGTTCTGTGGC	1739
Human IL-2 Probe	TCACTTCCTACCTTCTTGGGCATGTAA	1740
Human IL-2 Probe	TCACTTCCTACCTTCTTGGGCATGTAAAAC	1741
Human IL-2 Probe	TCACTTCCTACCTTCGGCCATGTAA- C18ddC	1742
Human IL-2 Invader	GAAGATGTTTCAGTTCTGTGG- ddC	1743
Human IL-2 Probe	FI- ACTTCCTACTICATICCATICAAAAIC	1744
Human IL-2 Probe	ACTTCCTACTICATICCATICAAAATC - C18ddC	1745
Human IL-2 Invader	GAGTTTGGGATTCTTGTAATTAT-ddC	1746
Human IL-2 Probe	FI- CGTGTTCTGTGGCGTATCTTAATTCCATTCAAAATC	1747
Human IL-2 Probe	CGTGTTCTGTGGCGTATCTITAATTCCATTCAAAATC	1748
Human IL-2 Invader	GAGTTTGGGATTCTTGTAATTAT - ddC	1749
Human IL-2 Probe	FI- CGTGTTCTGTGGCGTATCTTAATTCCATTCAAAATCATCTG	1750
Human IL-2 Probe	CGTGTTCTGTGGCGTATCITAATTCCATTCAAATCATCTG	1751
Human IL-2 Probe	FI- CGTGTTCTGTGGCGTATC <u>TTAATTCCATTCAAAATCATC</u>	1752
Human IL-2 Probe	CGTGTTCTGTGGCGTATCTTAATTCCATTCAAAATCATC	1753
Human IL-2 Invader	GAGTTTGGGATTCTTGTAATTAT-ddC	1754
HUMAN B-ACTIN		ł
Human β-actin Probe	FI-TTCCTACICTIGAICTICATIGIGG	1755
Human β-actin Invader	CTCAGGAGGAGCAATGATCTT	1756
Human β-actin Invader	CTCAGGAGGAGCAATGAT	1757
Human β-actin Probe	FI-TCACTTCCTACICIGGGICATCITCICG -C18ddC	1758
Human β-actin Probe	TCACTTCCTACICIGGGICAICTICICG -C18ddC	1759
Human β-actin Invader	GTGTTGAAGGTCTCAAACATGAT- ddC	1760
Human B-actin Invader	GGGTGTTGAAGGTCTCAAACATGAT - ddC	1761
Human β-actin Probe	FI- CGTGTTCTGTGGCGTATCTGGGTCATCTTCTCG	1762
Human β-actin Probe	CGTGTTCTGTGGCGTAICTGGGTCAICTICTCG	1763
Human β-actin Invader	GGGTGTTGAAGGTCTCAAACATGAT - ddC	1764
GAPDH Limon CABAL Broke	では、これでは、これでは、これでは、これでは、これでは、これでは、これでは、これ	
	TTCATACGGTTGGTTGACGTCAATG	1765
Human GAPDH Invader	GGAATCATATTGGAACATGTAAACCATC	1767
Human GAPDH Probe	FI- TTCATACGGTTGGCTCCTGGAAGATG	1768

FIGURE 47AP

Human GAPDH Probe Human GAPDH Invader Human/Mouse/Rat GAPDH Probe Mouse GAPDH Invader Mouse GAPDH Invader Mouse GAPDH Invader	TTCATACGGTTGGCTCCTGGAAGATG CACTTGATTTTGGAGGGATCTCA TTCATACGGTTGGTGGGTCAATG AGAATCATACTGGAACATGTAGACCATC FI-TGGCGTATCATGGAACATGTAGACCATC FI-TGGCGTATCATGGACTTGA GGAGTCATACTGGAACATGTAGACC TGGCGTATCATGGAACATGTAGACC TGGCGTATCATGGAACATGTAGACC GGAGTCATACTGGAACATGTAGACA GGAGTCATACTGGAACATGTAGACA	1769 1770 1771 1772 1773 1774 1775 1776
MOUSE IL-6 Mouse IL-6 Probe Mouse IL-6 Probe Mouse IL-6 Invader	FI- TGGCGTATC <u>ICITITCICATI</u> TGGCGTATC <u>ICITITCICATI</u> ACAATCAGAATTGCCATTGCACAACA	1779 1780 1781
MOUSE Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Invader	EI-GAAGGCAGAGCGTGAGGC GAAGGCAGAGGACCGTGAGGC AAGACATCTGGTGTTGTAGTGA	1782 1783 1784
Mouse Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Invader	FI-TGGCGTATC <u>TCCCCAGAGAAGC</u> TGGCGTATC <u>TCCCCAGAGAAGC</u> TGGCGTATC <u>TCCCCAGAGAAGC</u> CACTGAGCCGATGAAGGTGGTAA	1785 1786 1787
Mouse Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Invader	FI- TGGCGTATC <u>TAGGGCTCCAAGAG</u> TGGCGTATC <u>TAGGGCTCCAAGAG</u> GTGTTCAGGTTTTGGAGGCGCGATAA	1788 1789 1790
Mouse Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Invader FRET Probe	FI-TGGCGTATC <u>TAGGGCTCCAAG</u> TGGCGTATC <u>TAGGGCTCCAAG</u> GTGTTCAGGTTTTGGAGGCGGATAA FI-ATTC(CY3)TCTCAGA-3'NH2 FI-ATTC(CY3)TCTCAGAC-3'NH2	1791 1792 1793 1794
FRET Probe SRT Mouse Oncostatin M Arrestor Mouse Oncostatin M Arrestor Mouse Oncostatin M Arrestor	FI-ATTC(CY3)TCTCAGACT-3'NH2 CAGTCTGAGATGAATGG <u>CCAGG-</u> 3'NH2 CTTGGAGCCCTAGATA-NH2 CTTGGAGCCCTAGAT-NH2 CTTGGAGCCTAGA-NH2	1796 1797 1798 1799
Mouse Oncostatin M Probe Mouse Oncostatin M Probe Mouse Oncostatin M Invader SRT Arrestor	CTGGCGTATCI <u>AGGGCICCA</u> CCTGGCGTATC <u>IAGGGCICCA</u> GTGTTCAGGTTTTGGAGGCGATAA CAGTCTGAGATGATACG <u>CCAGG</u> -3'NH2 CTTGGAGCCTAGAI-NH2	1801 1802 1803 1804 1804
Mouse Oncostatin M Probe	FI-CTCTCTCTCTAGGGCTCCA	1806

FIGURE 47AQ

Mouse Oncostatin M Probe	CTCTCTCGTCTC <u>TAGGGCTCCA</u>	1807
Mouse Oncostatin M Invader	GTGTTCAGGTTTTGGAGGCGGATAA	1808
SK-	CAGTCTGAGATGAGACGAG <u>AGAGT</u> -NH2	1809
Mouse Oncostatin M Arrestor	CTTGGAGCCCTAGAG-NH2	1810
Mouse Oncostatin M Probe	FI- TGGCGTATCIAGGGCICCA	1811
Mouse Oncostatin M Probe	TGGCGTATCIAGGCCICCA	1812
Mouse Oncostatin M Invader	GTGTTCAGGTTTTGGAGGCGGATAA	1813
Mouse Oncostatin M Probe	TGGCGTATCICCCCAGAGAAA	1814
Mouse Oncostatin M Probe	TGGCGTATCICCCCAGAGA	1815
Mouse Oncostatin M Invader	CACTGAGCCGATGAAGCGATGGTAA	1816
Mouse Oncostatin M Probe	TGGCGTATCIAIAGGGCIC	1817
Mouse Oncostatin M Invader	GTGTGTTCAGGTTTTGGAGGCGGAA	1818
Mouse Oncostatin M Probe	CTCTCTCGTCTCTICAGGTTTTG	1819
Mouse Oncostatin M Invader	GGCAGCTCTCAGGTCAGGTGTGA	1820
Mouse Oncostatin M Invader	AGGCAGCTCTCAGGTCAGGTGTGA	1821
SRT	CAGTCTGAGATGAGACGAGAGAGI-NH2	1822
FRET Probe	FI-ATTC(CY3)TCTCAGAC-3'NH2	1823
Mouse Oncstatin M Arrestor	CAAAACCTGAAGAGA-3'NH2	1824
Mouse Oncostatin M Arrestor	CAAAACCTGAAGAG-3'NH2	1825
Mouse Oncostatin M Arrestor	CAAAACCIGAAGACG-3'NH2	1826
Mouse Oncostatin M Probe	FI- CTCTCCTCTTCAGGTTTTG	1827
Mouse Oncostatin M Probe	CTCTCTCGTCTTCAGGTTTTG-NH2	1828
Mouse Oncostatin M Invader	GGCAGCTCTCAGGTCAGGTGTGA	1829
Mouse Oncostatin M Stacker	GAGGCGGATATAGGGCT- Biotin TEG	1830
HUMANIONGOSITATIINIM		
Human Oncostatin M Probe	CTCTCTCGTCTTCTAAGGACTIA	1831
Human Oncostatin M Probe	CTCTCCTCGTCTTCIAAGGACTIAC	1832
Human Oncostatin M Invader	GAAACAGGAGTGCAAGGACCAGACA	1833
Human Oncostatin M Probe	TCACGTCTCTCAGGITITG	1834
Human Oncostatin M Probe	GTCACGTCTCTICAGGTTTTG	1835
Human Oncostatin M Probe	AGTCACGTCTC <u>TTCAGGTTTTG</u>	1836
Human Oncostatin M Probe	CAGTCACGTCTCTTCAGGTTTTG	1837
Human Oncostatin M Invader	AGGCAGCTCTCAGGTCAGGTGTGA	1838
Fret Probe 1	FI- CAAC(CY3)GCTTCCTCCG	1839
SRT	CGGAGGAAGCAGTTGGAGACGTGACTG <u>IGG</u> -NH2	1840
SRT with mismatch	CGGAAGAGCAGTTGGAGACGTGACTG <u>TGG</u> -NH2	1841
SRT with mismatch	CGGACGAAGCAGTTGGACGTGACTGTGG-NH2	1842

FIGURE 47AR

bold indicates 2' o-methyl bases

Oligo Type	Oligo Sequence	Oligo # SE(SEQ ID NO
SECONDARY SYSTEM: SET 1 FRET probe 1 secondary target	5-F-CAAC(CY3)GCTTCCTCCG-3' 5- CGGAAGAAGCAGTTGGTGCGCCTC <u>GTIAA</u> -NH2	DB04001F6 649-10-01	1843 1844
SET 2 FRET probe 1 secondary target	5'-F-CAAC(CY3)GCTTCCTCCG-3' 5'-CGGAAGAAGCAGTTGGAGGCGTGACGGT-NH2-3'	DB04001F6 641-60-03	1845 1846
h2C19 designs 2 probe stacker invader arrestor SET 1	5-AACGAGGCGCACGATGTCCATCGA-NH2-3' 5-TTCTTGGTGTTCTTTACTTTCTC-3' 5-GCAATCAATAAAGTCCCGAGGGTTGTTC 5'-TCGATGGACATCGTGCGC-3'	971-26-09 971-26-12 971-26-11 971-26-10	1847 1848 1849 1850
h 2D6 p450 designs probe stacker invader arrestor SET 2	5'-CCGTCACGCCTCTCACCCATCT-NH2-3' 5'-CTGGTCGCCGCACCT-3' 5'-TGTAGGGCATGTGAGCCTGGA-3' 5'-AGATGGGAGAGAGGCG-3'	971-11-01 971-11-04 971-11-03 971-11-02	1851 1852 1853 1854
probe stacker invader arrestor SET 2	5'-CCGTCACGCCTCGAAGCCCTGT-NH2-3' 5'-ACTTCGATGTCACGGGATGTCATAGG-3' 5'-GAGTGTCGTTCCCTTAGGGATGCGC-3' 5'-ACAGGGCTTCGAGGCG-3'	971-11-05 971-11-08 971-11-06	1855 1856 1857 1858
probe stacker invader arrestor SET 2	5'-CCGTCACGCCTCCCTGCTGAGAAG-NH2-3' 5'-GCAGGAAGGCCTCCG-3' 5'-CCCGAGGCATGCACGCGGGA-3' 5'-CTTTCTCAGCAGGGAGGCG-3'	971-11-09 971-11-12 971-11-11	1859 1860 1861 1862

FIGURE 47AS

CCGTCACG CCGTCACG CCGTCACG GCCAGGA TTTCTCAGA TTTCTCAGA TTCTCAGGA TTCTCAGGA TCTCAGGA CCGTCACG	5-CCGTCACGCCTCCCTGCTGAGAAA-HEX-3' 1051-12-06 1863 5-CCGTCACGCCTCCCTGCTGAGAAA-3' 1864 5-CCGTCACGCCTCCCTGCTGAGAAA-NH2-3' 971-38-01 1865 5-CCGAGGCATGCCACGCGGA-3' 971-38-03 1866 5-CCGAGGAAGGCCTCC-3' 971-38-03 1867 5-CCGAGGAAGGCCTCC-3' 971-38-02 1868	5'-CCGTCACGCCTCCCTGCTGAGA-NH2-3' 971-38-07 1869 971-11-11 5'-AAGGCAGGAAGGCCTCC-3' 971-38-09 171-38-08 1871	CCTCCCTGCTGAGAA-NH2-3' 971-38-04 1872 AGGCCTGG-3' 971-38-06 1873 AGGCCTGG-3' 971-38-05 1874 CCTCCCTGCTGAGAAAG-NH2-3' 971-11-09 1875	971-11-12 971-11-10 971-11-10 1051-48-01 971-01-03 971-01-04 1051-48-02	CCTCCACCATATCCC-HEX-3' 1051-12-02 1882 CCTCCACCATATCCC-3' 1051-12-01 1883 CCTCCACCATATCCC-NH2-3' 971-01-01 1884
		5-CCG I CACGCC I CCC I GAGA-NH2-3' 5'-AAGGCAGGAAGGCC I CC-3' 5'-ICTCAGCAGGGAGGCG-3'	5'-CCGTCACGCCTCCCTGCTGAGAA-NH2-3' 5'-AGGCAGGAAGGCCTGG-3' 5'-TTCTCAGCAGGGAGGCG-3' 5'-CCGTCACGCCTCCCTGCTGAGAAAG-NH2-3'	5'-GCAGGAAGGCCTCCG-3' 5'-CTTTCTCAGCAGGGAGGCG-3' 5'-AACGAGGCGCACCATATCCC-NH2-3' 5'-CCAGCGGTTTCCATTGGCAAAGATCAA-3' 5'-CGGAAGAATGGGTCGACCATG-3' 5'-GGGATATGGTGGCO-3'	5'-CCGTCACGCCTCCACCATATCCC-HEX-3' 5'-CCGTCACGCCTCCACATATCCC-3' 5'-CCGTCACGCCTCCACATATCCC-NH2-3' 5'-GGGATATGGTGGAGGCG-3'

	1886 1887 1888 1889	1890	1891	1892 1893 1894 1895	1896	1897	1898 1899 1900	1902	1903
	1051-48-03 971-01-10 971-01-11 1051-48-04	971-01-08 971-01-10 971-01-11	971-01-09	1051-48-05 1051-48-10 1051-48-06	1051-48-07 1051-48-10 1051-48-09	1051-48-08	1051-48-11 1051-48-16 1051-48-15 1051-48-12	1051-48-13 1051-48-16 1051-48-15	1051-48-14
FIGURE 47AT	5'-AACGAGGCGCACAGAGCTGATGAG-NH2-3' 5'-GAGAAGAGCTCAAACAGCTGGCCGAATAA-3' 5'-TGAAAAGTCTGGTAGAACAAGTTCAGC-3' 5'-CTCATCAGCTCTGGTGCG-3'	5'-CCGTCACGCCTCCAGAGCTGATGAG-NH2-3'	5'-CTCATCAGCTCTGGAGGCG-3'	5'-AACGAGGCGCACCCTTGGATTTC-NH2-3' 5'-CTGTTCAATCTCCCTGTAGACTCTCTA-3' 5'-CGAAGCTCCTCTATCAG-3' 5'-GAAATCCAAGGGTGCGC-3'	5-CCGTCACGCCTCCCTTGGATTTC-NH2-3'	5'-GAAATCCAAGGGAGGCG-3'	5'-AACGAGGCGCACTGAGGGCC-NH2-3' 5'-GGAAGAGGAGGTGGGGTCCAA-3' 5'-CCCTTGGATTTCCGAAG-3' 5'-GGCCCTCAGTGCGC-3'	5'-CCGTCACGCCTCTGAGGGCC-NH2-3'	5'-GGCCCTCAGAGGCG-3'
SET 2	probe invader stacker arrestor SET 1	probe	SET 2	h 2B6 p450 alt.splice designs2 p l s s a SET 1	عــ «	a SET2	p s a SET 1	a – 0	a SET 2

h2B6 p450 alt. Splice designs4

FIGURE 47AU

probe invader stacker arrestor SET 1	5'-AACGAGGCGCACAATACAGAGCTG-NH2-3' 5'-GAGAAGAGCTCAAACAGCTGGCCGC-3' 5'-ATGAGTGAAAAGTCTGGTAGAAC-3' 5'-CAGCTCTGTATTGTGCGC-3'	1051-48-17 1051-48-22 1051-48-21 1051-48-18	1904 1905 1906 1907
probe invader etacker	5'-CCGTCACGCCTCAATACAGAGCTG-NH2-3'	1051-48-19 1051-48-22	1908
arrestor SET 2	5'-CAGCTCTGTATTGAGGCG-3'	1051-48-20	1909
probe invader stacker arrestor SET 1	5-AACGAGGCGCACGGTTGAGGTTCTG-NH2-3' 5-CAGCAAAGAAGAGCGAGAGCGTGTTGAC-3' 5'-GTGGCTGAATTCACTGTG-3' 5'-CAGAACCTCAACCGTGCGC-3'	1051-48-23 1051-48-28 1051-48-27 1051-48-24	1910 1911 1912
probe invader stacker	5'-CCGTCACGCCTCGGTTGAGGTTCTG-NH2-3'	1051-48-25 1051-48-28 1051-48-27	1914
arrestor SET 2	5'-CAGAACCTCAACCGAGGCG-3'	1051-48-26	1915
h2B6 p450 designs probe invader stacker stacker arrestor SET 2	5'-CCGTCACGCCTCCACCATATCCCCG-NH2-3' 5'-CCGTCACGCCTCCACATATCCC-NH2-3' 5'-CGGAAGAATGGGTCGAC-3' 5'-CGGAAGAATGGGTCGACCATG-3' 5'-CGGAAGAATGGGTCGACCATG-3' 5'-GGGATATGGTGGAGGCG-3'	971-01-06 971-01-03 971-01-05 971-01-04	1916 1917 1918 1919
probe invader arrestor SET 2	5'-CCAGCGGTTTCCATTGGCAAAGATCAA-3' 5'-CGGGGATATGGTGGAGGCG-3'	971-01-01 971-01-03 971-01-07	1921
probe invader stacker	5'-CCGTCACGCCTCCAGAGCTGATGAG-NH2-3' 5'-GAGAAGAGCTCAAACAGCTGGCCGAATAA-3' 5'-TGAAAAAGTCTGGTAGAACAAGTTCAGC-3'	971-01-08 971-01-10 971-01-11	1923 1924 1925

FIGURE 47AV

arrestor SET 2	s'-ctcatcagctctggagggg-3'	971-01-09	1926
h2b6p450 designs 2 probe invader stacker arrestor SET 2	5'-CCGTCACGCCTCAGATGACTGCC-NH2-3' 5'-GGAGAAGGTCGGAAATCTCTGAATCTCATC-3' 5'-TCTGTGTATGGCATTTTGGCTCGG-3' 5'-GGCAGTCATCTGAGGCG-3'	971-01-12 971-01-13 971-01-15	1927 1928 1929 1930
h 2C19 designs 1 probe invader stacker arrestor SET 2	5'-CCGTCACGCCTCCATCCTTAATATCTAT-NH2-3' 5'-GAGAGATTGGTTAAGGATTTGCTGAA-3' 5'-CTGTAGGATATTTCCAATCACTGGG-3' 5'-ATAGATATTAAGGATGGAGGCG-3'	971-26-01 971-26-03 971-26-04 971-26-02	1931 1932 1933 1934
probe invader stacker arrestor SET 1	5'-AACGAGGCGCACCGTTCCAGGC-NH2-3' 5'-CATATCCATGCAGCACCACGATGA-3' 5'-CAAAATACAGAGTGAACACAGGGCC-3' 5'-GCCTGGAACGGTGCGC-3'	971-26-05 971-26-07 971-26-08 971-26-06	1935 1936 1937 1938
h2C19 shorter site 2 designs probe invader stacker arrestor SET 1	5'-AACGAGGCGCACCGTTCCAGG-NH2-3' 5'-CATATCCATGCAGCACCATGA-3' 5'-CCAAAATACAGAGTGAACACAGGGCC-3' 5'-CCTGGAACGGTGCGC-3'	971-68-01 971-26-07 971-68-03 971-68-02	1939 1940 1941 1942
probe probe probe invader stacker arrestor SET 1	5'-AACGAGGCGCACCGTTCCAGGC-NH2-3' 5'-AACGAGGCGCACCGTTCCAGGC-3' 5'-AACGAGGCGCACCGTTCCAGGC-HEX-3' 5'-CAAAATACAGAGTGAACACAGGGCC-3' 5'-GCCTGGAACGGGCC-3'	971-26-05 1051-12-03 1051-12-04 971-26-07 971-26-05	1943 1944 1945 1946 1947
rat 1A1, rat 1A2 probe	Rat 1A1 site 1 bs. 639-700 5'-CCGTCACGCCTCAGATTGACTATGCTG-NH2-3'	500-58-01	1948

FIGURE 47AW

invader stacker	5'-CAGTAACCTCCCCAAACTCATTGCTTC-3' 5'-AGCAGCTCTTGGTCATCGT-3'	500-58-03 500-58-04	1949 1950
arrestor SET 2	5'-CAGCATAGTCAATCTGAGGCG-3'	500-58-02	1951
rat 1A2 probe	Rat 1A2 site 1 bs. 674-725 5'-AACGAGGCGCACTGACATTCTCCAC-NH2-3'	500-58-05	1952
invader	5'-GTCCACAGCATTCCCTGAGGA-3'	500-58-07	1953
stacker arrestor	5'-AAAGICCTIGCTGCTCTTC-3' 5'-GTGGAGAATGTCAGTGCGC-3'	500-58-08 500-53-06	1954 1955
SET 1			
rat 2B1-2B2 patent			
probe	5'-AACGAGGCGCACTGGCTTGACACA-NH2-3'	500-49-05	1956
invader	5'-GICAATGTCCTTGGGAGCCAAAA-3'		1957
stacker	5'-GAGAAGTTCTGGAGGATGGTGG-3'	r2B1, 2B2 500-49-07	1958
arrestor SET 1	5'-TGTGTCAAGCCAGTGCGC-3'	500-49-06	1959
probe	5'-AACGAGGCGCACTGGCTTGACACAG-NH2-3'	500-49-01	1960
stacker	5'-AGAAGTTCTGGAGGATGGTGG-3'	300-49-03 r2B1 2B2 500-49-04	1961
arrestor	5'-CTGTGTCAAGCCAGTGCGC-3'	500-49-02	1962
SET 1			
rat 2B1-2B2 site 4	PROBE SET 2 (r2B1 bs 1299-1353, r2B2 bs. 474-528)		
probe	5'-AACGAGGCGCACGAGGAACAATTCATTT-NH2-3'	500-49-12	1963
invader	5'-GTTCTGGAGGATGGTGGTGAAGAAC-3'	500-49-10	1964
stacker	5'-CGGGCAATGCCTTCG-3'	500-49-14	1965
arrestor SET 2	5'-AAATGAATTGTTCCTCGTGCGC-3'	500-49-13	1966
probe	5-AACGAGGCGCACGAGGAACAATTCATTTC-NH2-3'	500-49-08	1967
invader		500-49-10	
stacker	5'-GGGCAATGCCTTCG-3'	500-49-11	1968
arrestor SET 1	5'-GAAATGAATTGTTCCTCGTGCGC-3'	500-49-09	1969
rat 2B1-2B2 ,5 patent			
probe	5'-AACGAGGCGCACAGCTGAGAAGCAG-NH2-3'	500-49-15	1970

FIGURE 47AX

1971 1972 1973 1974	1976 1977 1978 1979	1980	1982 1983 1984 1985	1986	1988 1989 1990
r2B1, 500-49-17 r2B2, 500-49-18 r2B1 500-49-20 r2B2 500-49-21 500-49-16	500-40-04 500-40-02 500-40-05 500-40-06	500-40-01 500-40-02 500-40-05 500-40-03	500-40-10 500-40-08 500-40-11 500-40-12	500-40-07 500-40-08 500-40-11 500-40-09	1073-19-06 500-40-14 500-40-17 500-40-15
5'-GCCTCAGCCGGATCACCGC-3' 5'-GCCTCAGCCCGATCACCGC-3' 5'-ATCTGGTACGTTGAGGTATT-3' 5'-ATCTGGTATGTTGGAGGTATT-3' 5'-CTGCTTCTCAGCTCTGCGC-3' Ire designed to detect both 2B1 and 2B2	Rat 2E1 PROBE SET (570C) 5'-CCGTCACGCCTCGTCGAAACGTTTGTT-NH2 5'-CCTCAGACACTTCTTGTCATTGTAC-3' 5'-CAAGAGGATATCCGCAATGACATTGC-3' 5'-AACAAACGTTTCGACGAGGCG-3'	5'-CCGTCACGCCTCGTCGAAACGTTTGTTGAAG-NH2-3' 5'-CTTCAACAAACGTTTCGACGAGGCG-3'	Rat 2E1 PROBE SET (822G) (designed over splice junction #5) 5'-CCGTCACGCCTCCTCTATG-NH2-3' 5'-GTTCTTGGCTGTTTTTCCTTA-3' 5'-AGGAGACAGTCACATC-3' 5'-CATAGAGATGGAGGGCG-3'	5'-CCGTCACGCCTCCTCTCTATGAG-NH2-3' 5'-CTCATAGAGATGGAGGGGGG-3'	Designed over splice junction #6 5'-CCGTCACGCCTCCTTCAATTTCTG-HEX-3' 5'-CCCTGTCAATTTCTTCATGAGGTTTA-3' 5'-CGTATTTCATGAGGATCAGGAGC-3' 5'-CCAGAAATTGAAGGAGGCG-3'
invader invader stacker stacker arrestor NOTE: all 3 invader/probe sets a	rat 2E1 p450 (afo61442) 500-73 p l s s SET 2	p I s SET 2	rat 2E1 p450 (afo61442) 500-73 p I s s a SET 2	p s s SET2	Rat 2E1 PROBE SET (969G) probe invader stacker arrestor SET 2

FIGURE 47AY

probe	5'-CCGTCACGCCTCCTCCATTTCTG-3'	1073-19-05	1992
probe probe invader stacker	5-CCGTCACGCCTCCTCCATTTCTGG-NH2	500-40-10 500-40-13 500-40-14 500-40-17	1993 1994
arrestor SET 2	5'-CAGAAATTGAAGAGGGGG-3'	500-40-18	1995
Rat 2E1 PROBE SET (969G) probe invader stacker	Designed over splice junction #6 5'-CCGTCACGCCTCCTCTCAATTTCT-NH2-3' 5'-CCCTGTCAATTTCTTCATGAAGTTTA-3' 5'-GGGTATTTCATGAGGATCAGGAG-3'	500-73-01 500-40-14 500-73-03	1996 1997 1998
arrestor SET 2	5'-AGAAATTGAAGAGGAGGCG-3'	500-73-02	1999
rat 3A's design 2 probe	5'-CCGTCACGCCTCGTTCCTGGGT-NH2-3'	500-43-15	2000
invader	5'-GAGCAAACCTCATGCCAATGCAC-3'	r3A1, 3A18 500-43-23	2001
invader	5'-GAGCAAACCTCATGTCAATGCAC-3'	r3A2 500-43-24	2002
invader	5'-GAGCAAACCTCATGCCAATACAC-3'	r3A2 500-43-24	2003
stacker	5-CCATTICCAAAGGGCAG-3' 5'-CCATTCCCAAGGGCAG-3'	short r3A1, 3A2, 3A18 500-43-19 short r3A9 500-43-20	2004
arrestor SET 2	5'-ACCCAGGAACGAGGCG-3'	500-43-16	2006
probe	5'-CCGTCACGCCTCGTTCCTGGGTC-NH2-3'	500-43-13	2007
invader		r3A1, 3A18 500-43-23	
invader arrestor SET 2	5'-GACCCAGGAACGAGGCG-3'	r3A2 500-43-24 500-43-14	2008
rat 3A's desing 3			
probe	5'-CCGTCACGCCTCTGAGAGCAAACCT-NH2-3'		2009
invader	5'-AGAGCGAGTTTCATATTCAA-3' 5'-AGAGCAACTTTCATGTTCAA-3'	r3A1, 3A2 500-43-35	2010
ropexei	S.ACAGCAAGTTTCATGTTGAA.3'	13/18 5/0-43-37	2017
stacker	5'-CATGCCAATGCAGTTCCTG-3'	r3A1. 3A18 500-43-31	2012
stacker	5'-CATGTCAATGCAGTTCCTG-3'	r3A2 500-43-32	2014
stacker	5'-CATGCCAATACAGTTCCTG-3'	r3A9 500-43-33	2015

FIGURE 47AZ

arrestor SET 2	5'-AGGTTTGCTCTCCGAGGCG-3'	500-43-30	2016
probe invader invader invader	5'-CCGTCACGCCTCTGAGAGCAAACCTCA-NH2-3'	500-43-27 r3A1, 3A2 500-43-35 r3A9 500-43-36 r3A18 500-43-37	2017
arrestor SET 2	5'-TGAGGTTTGCTCTCAGAGGCG-3'	500-43-28	2018
rat 3A's designs			
probe	5'-CCGTCACGCCTCGGAACATCTCCT-NH2-3'		2019
invader	5'-TGTCTCCATACTGTTCAATGATGGC-3'	r3A1, 3A2 500-43-09	2020
invader	5-IAICIGIAIACIGGI AAIGAIGGC-3	13A9 500-43-10	2021
	S-INICIONAL MCICIONI CARGO COSTA DE LA TONOTA DEL TONOTA DE LA TONOTA DEL TONOTA DE LA TONOTA DEL TONOTA DE LA TONOTA DE LA TONOTA DEL TONOTA DEL TONOTA DE LA TONOTA DE LA TONOTA DEL TONOTA DEL LA TONOTA DEL TONOT	284 282 500 42 06	2022
n u	S-TGAGICTTCCCACTGGTG-3	1341, 342 300-43-03	2023
, (5'-TGAGTTTGCCACTGGTG-3'	r3A18 500-43-07	202
SET 2			
probe	5'-CCGTCACGCCTCGGAACATCTCCTTGA-NH2-3'	500-43-01	2026
invader		r3A1, 3A2, 500-43-09)
invader		r3A9 500-43-10	
invader		r3A18 500-43-11	
arrestor	5'-TCAAGGAGATGTTCCGAGGCG-3'	500-43-02	2027
SET 2			
rat 3A's design 2b			
probe	5'-CCGTCACGCCTCGTTCCTGGG-NH2-3'	991-39-01	2028
invader	5'-GAGCAAACCTCATGCCAATGCAC-3'	r3A1, 3A18 500-43-23	2029
invader	5'-GAGCAAACCTCATGTCAATGCAC-3'		2030
invader	5'-GAGCAAACCTCATGCCAATACAC-3'	r3A9 500-43-25	2031
stacker	5' -TCCATTTCCAAAGGGCAG-3'	r3A1, 3A2, 3A18 991-39-03	2032
stacker	5' -TCCATTCCCAAAGGGCAG-3'	r3A9 991-39-04	2033
arrestor	5'-CCCAGGAACGAGGCG-3'	991-39-02	2034
SET 2			
rat or human 1A1 shorter site 2			
probe	5'-CCGTCACGCCTCCTGTGAT-HEX-3'	1073-19-02	2035
probe	5'-CCGTCACGCCTCCTGTGAT-3'	1073-19-01	2036

FIGURE 47BA

probe	5'-CCGTCACGCCTCCTGTCTGT-NH2-3' 5'-CCGTCACAATGCTCAATGAGGA-3'	991-12-04	2037
	5'-TCCTGACAGTGCTCAGGA-3' 5'-TCCTGACAGTGCTCAATCAGGA-3' 5'-GTCCCGGATGTGGCCC-3' 5'-ACATCACAGGAGGCG-3'	1 141 500-53-11 1 h 141 500-53-12 rat/human 141 991-12-06 500-53-10	2039 2039 2040 2041
	5'-CCGTCACGCCTCCTGTGTGATG-NH2-3'	991-12-01 r 1A1 500-53-11	2042
invader stacker arrestor SET 2	5'-TCCCGGATGTGGCCCT-3' 5'-CATCACAGACAGGGCG-3'	n 1A1 500-53-12 rat/human 1A1 991-12-03 991-12-02	2043 2044
	5'-CCGTCACGCCTCCTGTCTGATGT-NH2-3'	500-53-09 F 1A1 500-53-11	2045
invader stacker arrestor SET 2	5'-GTCCCGGATGTGGCCC-3' 5'-ATCACAGACAGGAGGCG-3'	h 1A1 500-53-12 rat/human 1A1 991-12-06 991-12-05	2046 2047
rat or human 1A1 site 1 probe invader stacker stacker arrestor	5'-CCGTCACGCCTCTGGCCCTTC-NH2-3' 5'-CTGTCTGTGATGTCCCGGATGA-3' 5'-TCAAATGTCCTGTAGTGCTC-3' 5'-TCAAAGGTTTTGTAGTGCTC-3' 5'-GAAGGGCCAGAGGCG-3'	500-53-04 500-53-03 rat 1A1 500-53-06 human 1A1 500-53-07 500-53-05	2048 2049 2050 2051 2051
SE 1.2 probe invader arrestor SET 2.	5'-CCGTCACGCCTCTGGCCCTTCTC-NH2-3' 5'-GAGAAGGGCCAGAGGGG-3'	500-53-01 500-53-03 500-53-02	2053
Rat/Human 1A1 site 2 probe invader invader stacker arrestor	5'-CCGTCACGCCTCCTGTGTGATGT-NH2-3' 5'-TCCTGACAATGCTCAATGAGGA-3' 5'-TCCTGACAGTGCTCAATCAGGA-3' 5'-CCCGGATGTGGCCCT-3' 5'-ACATCACAGAGAGGCG-3'	500-53-09 r 1A1 500-53-11 h 1A1 500-53-12 rat/human 1A1 500-53-14 500-53-10	2055 2056 2057 2058 2059

FIGURE 47BB

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rat or human 1A2 sites			
probe	5'-AACGAGGCGCACGGACTGTTTTCTGC-HEX-3'	1073-19-04	2060
probe	5'-AACGAGGCGCACGGACTGTTTTCTGC-3'	1073-19-03	2061
probe	5'-AACGAGGCGCACGGACTGTTTTCTGC-NH2-3'	500-53-15	2062
invader	5'-CTTGTTGAAGTCTTGATAGTGTTCCTC-3'	rat 1A2 500-53-17	2063
invader	5'-CTTGTCAAAGTCCTGATAGTGCTCCTC-3'	human 1A2 500-53-18	2064
arrestor	5'-GCAGAAACAGTCCGTGCGC-3'	500-53-16	2065
SET 1			
shorter h2C19 design site 3			
probe	5'-AACGAGGCGCACGATGTCCATCG-NH2-3'	971-48-01	2066
invader	5'-GCAATCAATAAAGTCCCGAGGGTTGTTC-3'	971-26-11	2067
stacker	5'-ATTCTTGGTGTTCTTTTACTTTCTC-3'	971-48-03	2068
arrestor	5'-CGATGGACATCGTGCGC-3'	971-48-02	2069
SE1 1			

FIGURE 47BC

			FIGURE 4/BC	/BC	
Human IL-10	IL-10				
Oligo Type	Sequence	Oligo Number	Secondary Cassette	Comments	OF OIL OF
probe	aacgaggcgcaccaaactcactcatggct-NH2	511-31-01	FV-1 & FV-2	3' amine	2070
arrestor	agocatgagtttggttgcg	511-31-02		All 2'-Ome + 3' amine arrestor for 511-31-01	2071
probe	aacgaggcgcaccaaactcactcatggc-NH2	511-30-01	FV-1 & FV-2	3' amine	202
arrestor	gccatgagtttggtgcg	511-30-02		All 2'-Ome + 3' amine arrestor for 511-30-01	2073
arrestor	gocatgagtttgg	380-89-02		All 2-Ome Same as 380-82-02	2074
arrestor	gocatgagtttggtg	380-89-04		All 2-Ome Same as 380-82-04	2075
arrestor	gocatgagtttggtgcg	380-89-08		All 2-Ome Same as 380-82-06	2076
arrestor	gocatgagtttggtgcgcc	380-89-08		All 2-0me Same as 380-82-08	2077
probe	aacgaggcgcaccaaactcactcatgg-NH2	511-67-01	FV-1 & FV-2	3'amine	2078
stacker	ctttgtacatgccttctcttggagc	781-79-01		stacker for 511-67-01 All 2'Ome	2079
arrestor	ocatgagtgagtttggtgcg	781-79-02		all 2'Ome arrestor for 511-67-01	2080
probe	aacgaggcgcaccaaactcactcatg-NH2	781-80-01	FV-1 & FV-2	3'amine	2081
stacker	gctttgtacatgccttctttggag	781-80-02		stacker for 781-80-01 All 2'Ome	2082
arrestor	catgagttgggtgcg	781-80-03		all 2'Ome arrestor for 781-80-01	2083
probe	aacgaggcgcaccaaactcactcat-NH2	781-81-01	FV-1 & FV-2	3'amine	2084
stacker	ggctttgtacatgccttctcttgga	781-81-02		stacker for 781-81-01 All 2'Ome	2085
stacker	ggctttgtagatgcctttctcttgga	938-74-01		stacker for 781-81-01 All 2'Ome to replace 781-81-02	2086
arrestor	atgagtgagtttggtgcg	781-81-03		all 2'Ome arrestor for 781-81-01	2087
probe	cogtcacgcctccaaactcactcat-NH2	938-46-02	MO4-1/MO4-2/MO4-3	same as 938-46-01 w/ 3' amine	2088
arrestor	atgagtgagtttggaggc	938-46-03		all 2'Ome arrestor for 938-46-01&02	2089
invader	taggcttctatgtagttgatgaagatgta	380-59-02			2090
invader	gtcatgtaggcttctatgtagttgatgaagatgta	511-32-01		longer invader 380-59-02	2091
Mouse IL-4	L4				
Oligo Type	Sequence	Oligo Number	Secondary Cassette	Comments	
probe	aacgaggcgcactctctgtgacctcg	511-14-01	FV-1 & FV-2		2092
arrestor	cgaggtcacaggagtgcg	511-14-02		All 2'-Ome + 3' amine arrestor for 511-14-01	2093
probe	aacgaggcgcactctcctgtgacct-NH2	511-12-01	FV-1 & FV-2	458-34-01 with 3' amine	2094
arrestor	aggicacaggagagtgcg	511-02-01		All 2'-Ome + 3' amine arrestor for 458-34-01	2095
probe	cagtcacgtctctctctgtgacct-NH2	511-16-01	MO2	3' amine	2096
arrestor	aggicacaggagagacg	511-16-02		All 2'-Ome + 3' amine arrestor for 511-16-01	2097
arrestor	aggtcacaggagagac	511-50-01		All 2'-Ome + 3' amine arrestor for 511-16-01	2098
probe	aaccagtcgtacgtctcctgtgacct	458-35-01	MISC-1		2099
arrestor	aggicacaggagaacgtac	511-03-01		All 2'-Ome + 3' amine arrestor for 458-35-01	2100
egord.	ocagicgiacgiclocigigacci	458-35-02	MISC-1		2101
arrestor	aggicacaggagagaggg	511-04-01		All 2'-Ome + 3' amine arrestor for 458-36-01	2102
9000	ancounty and a second a second and a second	430-30-01	MISC-2		2103
arrestor	ontracacoactoco	511-13-02	2-4-1 & 1-4-2		2104
adoru	STIN-cototototototototototototototototototot	784-74-04	EV-1 & EV.2	, c.	200
stacker	octoontoapaatoconatoatoto	781-71-02	2-1 8 1 1-2	5 armine 5x 784.74.04	2007
amestor	tracapagapathoon	781-71-02		All 2. Omo assessor for 784 74 04	700
Invader	atoxatchocatocotoxata	380-32-01			2108
Invader	atocatctcogtgaatggcgtcocta	380-32-02		Same as 380-32-01 but underlined base is mismatch to sequence	2110
		:	i		
probe	aacgaggcgcaccccttctcctgtgac-NH2	511-44-01	FV-1 & FV-2	3' amine	2111
anesio	90000000000000000000000000000000000000	511-44-02		All 4-Ume + 3 amine arrestor for 511-44-01	2112
arrector	acadadadadadadada	511.69.03	1 V-1 Q FV-2	All 9' Ome + 9' emine emerges for 644 69 04	2113
invader	ggcacatccatctccgtgcatggcgta	511-45-01			2115
•	•				
probe	cogtcacgcctcctgtgacctcgt-NH2	511-46-01	MO4-1/MO4-2/MO4-3	3' amine	2116

FIGURE 47BD

2117 2118 2119 2120 2121 2122 2123 2124 2125 2126	2127 2128 2129	2130 2132 2133 2133 2134	2135 2136 2137	2138 2139 2140	2141 2142 2143 2144 2145 2146 2146	2148 2149 2150 2151 2152	2153
All 2'-Ome + 3' amine arrestor for 511-46-01 3' amine All 2'-Ome + 3' amine arrestor for 511-69-01 3' amine All 2'-Ome stacker for 781-68-01 All 2'-Ome arrestor for 781-68-01 3' amine All 2'-Ome arrestor for 781-69-01 All 2'-Ome arrestor for 781-69-01	3' amine All 2'-Ome + 3' amine arrestor for 511-17-01	3' amine All 2' Ome arrestor for 781-83-01 3' amine All 2' Ome arrestor for 781-82-01	3' amine All 2' Ome arrestor for 781-84-01	Comments 3' amine All 2'-Ome + 3' amine arrestor for 511-19-01	Comments 3' amine All 2'-Ome + 3' amine arrestor for 511-24-01 3' amine All 2'-Ome + 3' amine arrestor for 511-23-01 All 2'-Ome + 3' amine arrestor for 511-20-01 All 2'-Ome + 3' amine arrestor for 511-20-01	Comments 3' amine (based on 685-27-01-1 base shorter) All 2'-Ome + 3' amine arrestor for 511-77-01 3' amine (based on 685-27-01-2 bases shorter) All 2'-Ome + 3' amine arrestor for 511-78-01	Comments 3' amine (based on 685-21-01)
MO4-1/MO4-2/MO4-3 MO4-1/MO4-2/MO4-3 MO4-1/MO4-2/MO4-3	M02	TT-1/TT-2 MO4-1/MO4-2/MO4-3	MO4-1/MO4-2/MO4-3	Secondary Cassette MO2	Secondary Cassette MO2 MO2 MO2	Secondary Cassette TT-1/TT-2 TT-1/TT-2	Secondary Cassette MO4-1/MO4-2/MO4-3
511-48-02 511-69-01 511-68-02 781-68-03 781-68-03 781-69-03 781-69-02 781-69-03	511-17-01 511-17-02 511-18-01	781-83-01 781-83-02 781-82-01 781-82-02 781-82-03	781-84-01 781-84-02 781-84-03	Oligo Number 511-19-01 511-19-02 511-20-01	Ollgo Number 511-24-01 511-24-02 511-23-01 511-23-02 511-21-01 511-21-01	Oligo Number 511-77-01 511-77-02 511-78-01 511-78-02 685-28-01	Oligo Number 511-79-01
acqaggicacaggaggagc ccgicacgccicctcigigaccic.NH2 gaggicacaggaggagc ccgicacgccicctcigigacc.NH2 toggicacaggagcacchH2 toggitcacaggaggagcacchcicacaggagcacchcicacaggagcacchciccigigac.NH2 ccgicacgccicctcigigac.NH2 ccgitcacaaga	cagtoaogictococtictocx-NH2 aggagaagggagacg gcacatocaticogigoatggoga	oogoogagateactoclqligaco-NH2 ggtoacaggagtgatc oogtcaogocctoclgtgaco-NH2 oogtgaatggogtcocttca ggtoacaggagaggog	oogloaegoctooctgtgaco-NH2 ogtgoatggogtoocttcta ggtcacagggaggog	L-2 Sequence captcacgicctiagittacaacagitactc.·NH2 agaglaactigtigaaaactaaagagacg gcactcaaatigtigticagagccca	FN-y Sequence Sequence capticacyctccttttgccapttcc-NH2 ggaactggccaaaggagagacg capticacyctccttttgccapttc-NH2 ggaactggcaaaggagagacg capticacyctccttttgccapttc-NH2 gaactggcaaaggagagacg capticacyctctccttttgccaptNH2 aactggcaaaaggagagacg gctctcgcaggatttcatgtcaccaa	TNF-cx Sequence ccgccggagatcactcgactgcctg-NH2 caggccagtcagatcactcgactccgg ccgccggagatcactcgactcccg ccgccggagatcactcgactccNH2 aggccagtcagatctcgg cct gtc act cgg ggt tcg aga aga tga a	IL-1β Sequence googtcacgcotctcatctgtttagggcc-NH2
probe arrestor probe stacker arrestor probe stacker arrestor probe stacker arrestor invader invader	probe arrestor invader	probe arrestor probe invader arrestor	probe invader arrestor	Mouse IL-2 Oligo Type Sec probe cag arrestor aga invader goa	Mouse IFN-y Olgo Type Seque probe cagic arrestor ggada arrestor agact probe cagic arrestor acact invader gctct	Human TNF-ca Oligo Type Sequent probe cagocaga arrestor cagogaga probe cagocaga arrestor aggocaga invader ctt gtc ac	Human IL-1β Oligo Type Seque probe googto

FIGURE 47BE

2154 2155 2156	2157 2158 2159 2160 2161 2162 2165 2165 2166 2167 2169	-	2172 2173	2174 2175	2176 2177	2178 2179	2180 2181	2182 2183	2184 2185	2186 2187	2188 2189	2190 2191
All 2-Ome + 3' amine arrestor for 511-79-01 All 2-Ome + 3' amine arrestor for 511-79-01	Comments 3 amine (based on 685-16-01) All 2-Ome + 3' amine arrestor for 511-81-01 All 2-Ome + 3' amine arrestor for 511-81-01 3 lamine (511-81-01 with new arm) All 2-Ome + 3' amine arrestor for 781-27-01 3' amine (based on 685-16-01) All 2-Ome + 3' amine arrestor for 781-27-01 All 2-Ome + 3' amine arrestor for 511-81-01 3' amine (511-83-01 with new arm) All 2-Ome + 3' amine arrestor for 511-81-01 3' amine (781-29-01 with new arm) All 2-Ome + 3' amine arrestor for 781-29-01 3' amine (781-29-01 with new arm) All 2-Ome + 3' amine arrestor for 781-29-01 3' amine (781-29-01 with new arm) All 2-Ome + 3' amine arrestor for 781-29-01											
	Secondary Cassette MO4-1/MO4-2/MO4-3 MO4-1/MO4-2/MO4-3 MO4-1/MO4-2/MO4-3 MO4-1/MO4-2/MO4-3 TT-1/TT-2		FV-1	图FV-2	MO4-1	MO4-2	MO4-3	Ē	П-2	MO2	MISC-1	MISC-2
511-80-01 511-80-02 685-23-01	Oligo Number 511-81-01 511-82-01 511-82-02 781-27-01 781-27-01 781-28-01 781-28-02 781-29-01 781-29-01 781-30-01 781-30-02		277-68-05 187-46-01	(995-29-01) FV-2 767-29-02	641-60-03 187-46-01	562-93-01 187-46-01	896-29-02 MAN MO4-3 767-29-02	562-92-01 187-46-01	685-56-01 187-46-01	491-68-02 491-68-01	458-35-03 187-46-01	441-31-02 187-46-01
ggooctaaacagatgagaggogt ggooctaaacagatgagaggogtga caggtoctggaaggagcacta	Fig. 6 Sequence googlaagoctictoctoatigaatoct-NH2 aggattcaatgaaggagaggagggoogt cogtoagoctictoctoatigaatoct-NH2 aggattcaatgaggagagaggoogt cogtoagoctictoctoatigaatoct-NH2 aggattcaatgaggagagagaggoogt googlaaggagagagaggogt cogtoagoctictoctoctattgaatoc-NH2 ggattcaatgaggagagaggogt cogtoagoctictoctoctattgaatoc-NH2 ggattcaatgaggagagaggogt cogtoagoctictoctoctattgaatoc-NH2 gattcaatgaggagagaggogt cogtoagoctictoctoctattgaatoc-NH2 gattcaatgaggagagagagog cogtoagoctictoctoctattgaatoc-NH2 gattcaatgaggagagagagog cogtoagoctictoctoctattgaatoc-NH2 gattcaatgaggagagagog cogtoagoctoctoctoctattgaatoc-NH2 gattcaatgaggagagagog cogtoagoctoctoctoctattgaatoc-NH2 gattcaatgaggagagagog cogtoagoctoctoctoctattgaatoc-NH2 gattcaatgaggagagagog cogtoagoctoctoctoctoctoctoctoctoctoctoctoctocto	Secondary Cassettes	oggaggaagoagttggtgogooto gttaaNH2 Foaac(Cy3)gottcotoog	ccaggaagcaagtgglgcgcctcgttt Fcac(Z21)igcttcgtgg	cggaagaagcagtgaggcgtgacggtNH2 Fcaac(Cy3)gcttcctccg	oggaagaagogttggaggogtgaog gc NH2 Fcaac(Cy3)gcttcctcog	ccaggaagcaagtggaggcgtgac ggu Fcac(221)tgcttcgtgg	oggaggaagcagttggtgatctoggo gg NH2 Fcaac(Cy3)gcttoctoog	oggaagaagcagttggtgatctoggooggNH2 Fcaac(Cy3)gcttcctoog	gctactgagatgaaggagacgtgactgtaNH2 Fcttc(Cy3)tctcagtagc	oog agg aag cgg ttg ogt aog act g <u>gt taa</u> -NH2 Fcaac(Cy3)gcttoctoog	ogg agg aag ogg ttg gtg ogg gtg gtt gg PO3 Foaao(Cy3)gottootoog
arrestor arrestor invader	Human II. Oligo Type probe arrestor probe	Seconda	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe	SRT FRET probe

FIGURE 47BF

Oligo sequence descriptions: 5' to 3' direction, 2'-Ome nts are bolded and underlined, internal modifications defined in ()

FRET Oligo/SRT Combinations

	ERET Office	Las			
Set 1	187-46-01	641-60-02			
Set 2	187-46-01	690-82-03			
Star	307-70-02	339-50-03			
A tag	303.18.05	343-63-07			
Set	303-18-05	343-25-01			
	187.46-01	640-40-04			
Set 7	744-80-03	277.080.08N			
- e	187-46-03	822-18-07			
0 10	767 20 63	124 14 40			
0000	20-07-101	01:1/-1/1			
04 10	70-67-797	10-67-986			
Set 11	10-07-7-901	996-29-01			
Set 12	307-70-02	307-70-04			
Set 13	491-01-01	491-02-04			
Set 14	187-46-01	562-84-01			
			•		
	# 00110	Ollo Seguence			ON CLOSE
	187-46-01	Eam_CAACICY3/GCTTCCTCCA			3103
	307-70-02	Fam-ATTC/CY3)TCTCAGAC-NH2			2192
	303-18-05	Fam-TAAC(CY3)GCTTCCTCCG			2100
	744-80-03	Fam-CAA(Dabox)TGCTTCCTCG			2105
	767-28-03	Red Dve-CTC(Z-21)TTCTCAGTGCG			2196
	767-29-02	Fam-CAC/Z-21)TGCTTCGTGG			2107
	1067-20-01	Fam-CAC/Z-28)TGCTTCGTGG			2108
	491-01-01	Fam-CTTC(CY3)TCTCAGAC			2199
		SRT			
	Oligo #	Oligo Sequence			SEQ ID NO
	641-60-02	CGGAGGAAGCAGTTGGAGGCGTGACGGT-NH2			2200
	690-82-03	CGGAGGAAGCAGTTGTGGCGGTGACGGTT			2201
	339-50-03	CAGTCTGAGATGAATGAGACGAGAGAGT-NH2			2202
	343-63-07	CGGAGGAAGCGGTTAGTCTGTCACGICAI-NH2			2203
	343-25-01	CGGAGGAAGCGGTTAGTCTGCCACGICAI-NH2			2204
	649-10-01	CGGAAGAAGCAGTTGGTGCGCCTCGTTAA-NH2			2205
	277-068-05N	CGGAGGAAGCAGTTGGTGCGCCTCGTTAA-NH2			2206
	833-18-07	CGGAGGAAGCAGTTGCGGCGTGCGGCI-NH2			2207
	777-71-10	GCGCAGTGAGAATGAGGAGGCGTGACGG <u>U</u> -NH2			2208
	996-29-01	CCAGGAAGCAAGTGGTGCGCCTCGUUU			2209
	307-70-04	CAGTCTGAGATGATGCTACGCCAGG-NH2			2210
	491-02-04 562-84-01	AGTCTGAGATGAAGGAGACGTGACTG <u>TGG</u> -NH2 CGGAGGAAGCGGTTGGTGATCTCGGCG-NH2			2211
					!
Oligo Type	Oligo #	Oligo Sequence	Notes	Position	SEQ ID NO
Human IL-2					
Probe	196-56-01	TCTGTGGCGTATCCTTGGGCATGTAA		Cacitor transfer 2	2213
Probe	196-56-02	GTGGCGTATCCTTGGGCATGTAA		t ilonomore Soudo	2214
Probe	196-56-03	GCGTATCCTTGGGCATGTAA			2215
Invader	128-93-02	GAAGATGTTTCAGTTCTGTGG(ddC)	ddC = dideoxy C		2216
Capture Oligo	145-030-05	AAAAGATACGCCACAGAACACG(BIOTIN-dA)TT			2217
Probe	315-28-01	TGGCGTATCTTAATTCCATTCAAAAT		Splice Junction 1	2218
Invader	315-28-02	TGGGAGTTTGGGATTCTTGTAATTAA			2219

FIGURE 47BG

2220 2221 2222 2223 2224 2225 2225 2226 2226 2228	2230 2231 2233 2234 2236 2240 2241 2241 2245 2245 2246 2246 2246 2246 2246 2246	2256 2257 2257 2258 2260 2261 2263 2264 2265 2265 2266 2266 2266 2266
Splice Junction 1 Splice Junction 1 Splice Junction 2	Splice Junction 5 Splice Junction 4 Splice Junction 4 Splice Junction 3 Splice Junction 3 Splice Junction 3 Splice Junction 3	Splice Junction 6 Splice Junction 6 Splice Junction 4 Splice Junction 4 Splice Junction 4
		FI = Fluorescien FI = Fluorescien FI = Fluorescien Same as 425-59-01 without Fluorescien
AAAGGTACGCCACGC(BIOTIN-dT)C TGGCGTATCTAATTATTAATTCCATTC ATCCTGGTGAGTTTGGGATTCTTGA AAAAGATACGCCACACGC(BIOTIN-dT)C TGGCGTATCTTCATTCAATCAATCATC GTTTGGGATTCTTGTAATTATTAAA AAAGATACGCCACACG(BIOTIN-dT)C GTGGCGTATCCTTGTGGGCAT GAAGATACCTCTTGTGGGCAT GAAGATACCTCTTGGGCAT GAAGATGCTTCTGTGGCCAT GAAGATGCTTCTGTGGCCAT	TGGCGTATCTCTGGGTCATCTTC GGGTGTTGAAGGTCTCAAACATGAA AAAGGATACGCCACAGC(BIOTIN-dT)C TGGCGTATCTCTTGATCTTCATTGT ACATGGGCTCAGGAGGAGCAATGAA AAAAGATACGCCACAGC(BIOTIN-dT)C TGGCGTATCTGATCTGGGTCATCT TGGCGTATCTGATCTG	FI-CTCTCGTCTCCTGGAAGA ATTTGATGTGGGGGTCTCGCA FI-CTCTCGTCGTGCTGACAATC GCAGTTGGTGGTGCAGGATGCATA FI-CTCTCGTCTCTCTCTCTGCAGAATG GCAGTTGTCTCTCTCTCTCTCTGCAA FI-CTCTCTCGTCTCTCTCTGCAA GATTGATGTTAGTGGGGTCTCGA CATTGATGTTAGTGGGGTCTCGA CTCTCTCGTCTTCTTGAA CTCTCTCGTCTCTCTGGAAG CTCTCTCGTCTCCTCGGAAG CTCTCTCGTCTCCTCGGAAG CTCTCTCGTCTCCTCGGAAG Set 3 CTCTCTCGTCCTCCTGGAAG CTCTCTCGTCCTCCTGGAAG CTCTCTCGTCCTCCTGGAAG CTCTCTCGTCTCCTCGGAAG CTCTCTCGTCGTCTCCTCGGAAG CTTTGATGTTAGTGGGGGTCTCGA
195-023-01 315-29-01 315-29-02 195-023-01 315-29-03 315-29-04 195-023-01 315-30-02	315-26-01 315-26-02 195-023-01 315-27-02 195-023-01 315-91-02 195-023-01 315-92-01 315-92-01 315-92-01 315-92-01 315-92-01 315-92-01 340-32-01 340-33-01 340-33-01 740-01-02 740-01-03 740-01-04 740-01-09	425-59-01 425-59-02 425-50-01 425-60-02 - 425-61-01 425-80-01 425-80-02 425-80-02 425-89-02 425-89-02 425-89-02
Capture Oligo Probe Invader Capture Oligo Probe Invader Capture Oligo Probe Invader Capture Oligo	Human b-actin Probe Invader Capture Oligo Probe Invader Sacondary Cassette Probe Invader Arrestor Secondary Cassette Stacker Invader Stacker Invader Sacondary Cassette	Mouse GAPDH Probe Invader Arrestor Secondary Cassette Probe Invader

FIGURE 47BH

425-87-05
425-87-03 CTCTCTCGTCTTACCAGGAAATG 425-61-02 GCTGTAGCCGTATTCATTGTCAA 425-87-06 <u>CATTCCTGGTAGAGAGG</u> 5-81
453-23-01 ATGACGTGACAGACCTCCTGGAAGAT 453-23-03 ATGACGTGACAGACCTCCTGGAAGATG 425-80-02 CAITTGATGTTAGTGGGGTCTGGA 59:23-04 Set 4
453-23-02 ATGACGTGGCAGACCTCCTGGAAGAT 425-80-02 CATTTGATGTTAGTGGGGTCTCGA 453-23-05 AICIICCAGGAGGICIGG-NH2 Set 5
524-51-05 TCGCTACTGAGATGAAGGAGACGTGACTGTA·NH2 524-51-06 TCGCTAATGAGATGAAGGAGACGTGACTGTA·NH2
796-72-01 AACGAGGCGCACCTITACATITICIATCGTATCC 428-81-02 CCITCCITATCCTGGAICITGGCA GAIS-2-02 GGAIACGAIAGAAAAIGIAAAGGIGCGC Set 6
796-72-03 AACGAGGCGCCTTTACCATTTCTATCGTATC 428-81-02 CCTTCCTTATCGTGGATCTTGGCA GAIACGAIGTAAAGGIGCGC 58-78-73-74-73-74-74-74-74-74-74-74-74-74-74-74-74-74-
820-35-01 AACGAGGCGCACCTTTACATTTCTATCG 820-35-02 AACGAGGCGCACCTTTACATTTCTATCGT 428-81-02 CCTTCCTTATCCTGGATCTTGGCA 820-35-03 ACGATAGAAAATGTAAAGGTGCGC 84 7 86 7
820-88-01 AACGAGGCGCACCTTTACATTTCTATCGT_UND 820-88-02 AACGAGGCGCACCTTTACATTTCTATCGT_U 820-88-03 AACGAGGCGCACCTTTACATTTTTATCGT_G
820-88-04 AACGAGGGCACCTTTACATTTCTATCGTT 428-81-02 CCTTCCTTATCCTGGATCTTGGCA 820-35-03 AGGATAGAAATGTAAAGGTGCGC
847-85-01 GCGGCACGCCCTTTACATTITCTATGGT 428-81-02 CCTTCCTTATCCTGGATCTTGGCA 847-85-02 AGGAIAGAAAAIGTAAAGGGGCG 847-85-03 AGGAIAGAAAAIGTAAAGGGGGGT
838-61-01 ACCAGGGGGGCGTTTACATTTCTATCGTATCCG

FIGURE 47BI

Arrestor Secondary Cassette	936-61-02	CGGATACGATAGAAATGTAAAGGTGCGC Set 7	Same as 428-87-03 without Biotin blocking group	2312
Monocyte Chemotactic Protein 1 (MCP-1) Probe Invader Arrestor Secondary Cassette	820-89-01 685-76-01 820-89-02	CCGTOACGCCTCCTTCGGAGTTTGGG GGGTTGTGGAGTGATGTTCAAGTA <u>CCCAAACTCCGAAGGAGGG</u> Set 9	Same as 720-92-01 without the amine	2313 2314 2315
MAGE-3 Probe Invader Stacker Probe Stacker Probe Invader Amschor	1001-01-01 871-18-03 871-18-01 1138-50-01 1138-50-02 1138-50-04 1138-50-05	FITTTOTGGAAGCTTTGCT CGATGCCAAAGACCAGCTGCAAGGAAG GAAGTCACAGGAAAATAC GCAGCTTGTTGGGA AACGAGGGGACGTTGGGTGA GCAGCTTCTTGGGACT AACGAGGGCGACGTTGGGTGA CCAGCGGCACGTTGGGTGA CTCCAGGCGACGTTGCGTGGCTGGC CTCCAGCCAACGTTGCTGCCTCCCCCACGCTTCCCCCCAACTCC CTCCACCCAACGTGCGCC CTCCACCCAACGTGCGCC CTCCACCCAACGTGCGCC CTCCACCCCAACGTGCGCC CTCCACCCCAACGTGCGCC CTCCACCCCAACGTGCGCC CTCCACCCCAACGTGCGC CTCCACCCCAACGTGCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCC CTCCACCCCAACCTCCCCC CTCCACCCCAACCTCCCCCACCCC CTCCACCCCAACCTCCCCCCCACCCC CTCCACCCCAACCTCCCCCCACCCC CTCCACCCCAACCTCCCCCCACCCCCACCCC CTCCCACCCCAACCTCCCCCCACCCCACCCCCCACCCCCACCCCCACCCCCC	Same analyte specific Region as 871-18-02.	2316 2317 2318 2319 2320 2321 2322 2323
Secondary Cassette Stacker Probe Stacker Probe Invader Arrestor	1138-51-01 1138-51-02 1138-51-03 1138-51-04 1138-51-05	Set 10 AGCITCTIGGGAIC AACGAGGCGCACTTGGGTGAGC GCITCTIGGGAICC AACGAGGCGCACTTGGGTGAGCA CAGGTACTTTCCTGCACAATGA IGCTCACCCAAGTGCGC		2325 2326 2327 2328 2329 2330
Secondary Cassette Stacker Probe Invader Arrestor Secondary Cassette Stacker Probe Invader Arrestor Secondary Cassette	1138-67-01 1138-67-02 1138-67-04 1138-67-06 1138-67-06 1138-67-06 1138-67-06	Set 11 TGGGGGTCGCTGCC ACGAGGGCCCACCATCATACA GGCCCTTGGACCCCAA IGTTATGAATTGGTGGTGCGC Set 11 ACGAGGATCACTGC ACGAGGATCACTGC ACGAGGCGCACCACTCATAA AGGGGCCCTTGGACCCACTTCATAA SGGCCTTGGACCCA Set 11 Set 11		2331 2332 2333 2334 2335 2336 2337 2338
Human Oncostatin M Probe invader Arrestor Arrestor Arrestor Arrestor Secondary Cassette Probe Invader Stacker Arrestor	339-30-02 284-42-03 374-32-01 374-32-02 374-32-03 524-39-01 435-40-02 389-47-07	CCTGGCGTATCTAGGGCTCCA GTGTTCAGGTTTTGGAGGCGGATAA CITGGAGCCTAGAIAG-NH2 CITGGAGCCTAGAIAGG-NH2 CTGGAGCCTAGAIAGG-NH2 Set 12 CAGTCACGTCTTCAGGTTTTG-NH2 AGGCAGCTTCAAGTCAGGTTTTG-NH2 AGGCAGGTTTCAAGTCAGGTCAA GAAAACCTGAAGAGGG-NH2	Same as 435-67-04 with 3' Amine	2339 2340 2341 2342 2342 2344 2344 2345 2346
Secondary Cassette Probe Arrestor Probe Probe Invader	1088-74-01 1088-74-02 1088-74-03 1088-74-04 603-75-03	Set 13 AACGAGGCCCTCTGTGTG CACAGAGGGTGCGC AACGAGGCACCCTTGTGTG-NH2 AACGAGGCCACCCTTGTGTG-NH2 AACGAGGCCACCCTTGTGTG-HEX GCAAGGACCAGACTGAGCGTA	HEX = Hexanediol	2348 2349 2350 2351 2352

FIGURE 47BJ

Stacker	752-01-05		2353
Secondary Consults	641-62-04	<u>CAGAGGGACG-</u> NH2	2354
Secondary Cassene	1100 1000		
Pioce States	1138-49-02	OTICI GGAG-NH2	2355
State	138-89-01		2356
Invader	1138-49-03	GrCTGA	2357
Arrestor	1138-49-04	NGAAGGTGCGC	2358
Secondary Cassette			
Probe	1138-49-06	AACGAGGCGCACTCTGCTTCT-NH2	2359
Stacker	1138-49-05		2360
Invader	1138-49-07	GAGATCTGA	2364
Arrestor	1138-49-08		2362
Secondary Cassette			7007
Probe	1138-40-10	A A A A A A A A A A A A A A A A A A A	0
Stacker	1130 40 00		2363
	1130-49-03		2364
roteon	1130 40 11		2365
Constant Constant	71-64-0011	ווארופעונענו	2366
Secondary Cassette			
Stacker	1163-01-01		2367
Probe	1163-01-02	NH2	2368
Invader	1163-01-03	TCCTGCATGAGATCTGTCTGCA	2369
Arrestor	1163-01-04	GCICCAGAAGIGCGC	2370
Secondary Cassette			· · ·
Stacker	1163-01-05	GGCCAAGGAGCAC	2371
Probe	1163-01-06	TGGAGCT-NH2	2372
Invader	1163-01-07		2373
Arrestor	1163-01-08		2374
Secondary Cassette			;
Stacker	1163-01-09	GCCAAGGAGCACG	2375
Probe	1163-01-10	GGAAGTC-NH2	22.0
Invader	1163-01-11		2377
Amestor	1183.01.12		727
Secondary Cassette	71-10-501		2378
84h6r			
Probe	688-51-01	CGCCGAGATCACGCCAACGACGACGTCT	2370
Invader	688-51-02	CACTA	2380
Arrestor	688-51-03		2381
Secondary Cassette			7007
Probe	688-51-04	SAGATCACCTCAACACCATAAAAGCCA	2382
Invader	688-51-05		2383
Arrestor	688-51-06		2007
Secondary Cassette			1007
ZHOM!			
7.00e	690-32-02		2385
Invader	690-32-04	GATCAAAGAGGC	2386
Stacker	709-52-01		2387
Arrestor	690-32-05	24GGCTTGAGGGGATC	2388
Secondary Cassette		Set 1	

FIGURE 47BK

bold indi	bold indicates 2' O methyl base	ıyl base		SEQ ID NO
ELISA Format K Leukocyte-associ G4731 Probe Set	ELISA Format Kits Leukocyte-associated mo G4731 Probe Set	ELISA Format Kits Leukocyte-associated molecule-1 alpha subunit, human (h-LFA1) G4731 Probe Set	uman (h-LFA1)	
a_			5'-CTCTCTCGTCTCCAGGGCGTCGTCGG-PO4-3' 5'-CTGTCACACGTCGGTGCTGA-3'	2389 2390
ပ			5'-AAAAAGGAGACAAGAGTG-3'	2391
for the rei	for the remainder of the oligo se	ligo sets on this list, the f	is on this list, the fret/target secondary sets are one of the following 11:	
FRET/TA	FRET/TARGET SETS	!		
set 1	FRET 307-70-03	TARGET 502-93-01		
set 2	307-70-03	502-93-02		
set 3	187-46-01	641-60-02		
set 4	187-46-01	277-68-05		
set 5	187-46-01	685-56-01		
set 6	187-46-01	641-60-03		
set 7	187-46-01	649-10-01		
set 8	680-17-02	782-70-02		
set 9	187-46-01	277-68-06		
set 10	187-46-01	491-02-02		
set 11	307-70-03	761-40-02		
FRETS				
004 400	c			C
187-46-01	o -		5'-Fam-CAAC (CY3);CTCCTCCG-3'	2392
680-17-02	. 2		5'-Fam-CGCT (CY3)TCTCGCTCGC-3'	2394
TARGETS	γ			
502-93-01	_		5'-CAGTCTGAGATGATGATACGAGAGAGT-NH2-3'	2395
502-93-02	2		5'-CAGTCTGAGATGAATGAGACGAGAGAGT-NH2-3'	2396
641-60-02	2		5'-CGGAGGAAGCAGTTGGAGGCGTGACGGT-NH2-3'	2397
277-68-05	5	-	5'-CGGAGGAAGCAGTTGGTGCGCCTCGTTAA-P04-3'	2398
685-56-01	Ξ		5'-GCGGAAGAAGCGGTTGGTGATCTCGGCGG-NH2-3'	2399
641-60-03	0		5'-CGGAAGAAGCAGTTGGAGGCGTGACGGT-NH2-3'	2400
649-10-01	Ξ		5'-CGGAAGAAGCAGTTGGTGCGCCTCGTTAA-NH2-3'	2401
782-70-02	2		5'-GCGAGAGAGACGCCCAAACCTGCCGTTC-3'	2402
277-68-06	90		5'-CGGAGGAAGCAGTTGTCCGCGAAGATG-3'	2403
491-02-02	2		5'-CGGAAGAAGCAGTTGGAGACGTGACTGTGG-NH2-3'	2404

FIGURE 47BL

761-40-02	5'-GGAGTGAGACAGCGAAAGACTGCCGTTCT-3'	2405
Cell Lysate Kits adipocyte lipid binding protein, mouse (m-aP2) C289 Probe Set p a a a a b p p p p p p p p p p a a a a	FRET/TARGET SET 1 5-CCGCCATCTAGGGTTATGATGCTA-3' 5-CCGCCATCTAGGGTTATGATGCTA-3' 5-CCGCCATCTAGGGTTATGATGCTA-3' 5-CTCTCTCGTCCTTCACCTTCCTGTCG-NH2-3' 3-P04-AGCAGGAAGTGGAAGGACAGC-5' 3-NH2-AGCAGCAGCAGCAGCG-5' 5-AACGAGCGCACCTTCACCTTCCTGTCG-NH2-3; 5-AACGAGCCCACCTTCACCTTCCTGTCG-NH1-3' 5-AACGAGCCCACCTTCACCTTCCTGTCG-NH1-3' 5-CATCTTCGCGGAAGTGGAAGGACAGC-5' 3-P04-GCCTGAAGTGGAAGGACAGC-5' 3-P04-GCCTGAAGTGGAAGGACAGC-5' 3-P04-GCCCTGAAGTGGAAGGACAGC-5' 5-CATCTTCCCCGTGCTTCCTGTCG-NH2 3-P04-GCCCTGAAGTGGAAGGACAGC-5' 5-CTTGCTCCCCGTGCTTCACCTTCCTGTCG-NH2 5-CTTGCTCCCCGTGCTTCACCTTCCTGTCG-NH2 5-CTTGCTCCCCGTGCTTCACCTTCCTGTCG-Biotin 3-P04-GGCCCCGAAGTGGAAGGACAGC-5' 3-P04-AGGGCCACCAAGTGGAAGGACGC-5'	2406 2407 2408 2408 2410 2411 2412 2413 2415 2415 2415 2419
G392 Probe Set p I	FRET/TARGET SET 1 5'-CTCTCTCGTCTCCACATTCCACCAG-NH2-3' 5'-TTGTGTAAGTCACGCCTTTCATAAT-3'	2422 2423
rev-ErbA, mouse (m-revErbA C155 Probe Set p	FRET/TARGET SET 4 5'-AACGAGGCGCACGAAGCAGGGTAATGAATCT-NH2-3' 5'-CCACTCCTGAAGGCTCCGCAGTC-3'	2424 2425
Carnitine palmitolytransferase, mouse (m-CPT-1) T352 Probe Set p I	FRET/TARGET SET 2 5-CTCTCTCGTCTCAATGCCTGTCGCC-NH2-3' 5-GCTTCAGGGTTTGTCGGAAGAAGAAC-3'	2426 2427
C851 Probe Set p .i	FRET/TARGET SET 2 5'-CTCTCTCGTCTCGTTTGCGGCGATACAT-NH2-3' 5'-CGGCTTGATCTCTTCACGGTCCAC-3'	2428 2429

Carnitine palmitolytransferase, human (h-CPT-1)

FIGURE 47BM

U744 Probe set p I a	FRET/TARGET SET 2 5'-CTCTCTCGTCTCAACTTCCAACTGTAATCT-NH2-3' 5'-CTCACGTAATTTGTAGCCCACCAGGAGTTTC-3' 3'-NH2-GCAGAGTTGAAGTTTATGGTGACATTAGA-5' 5'-TGGTCCAAGACCGACAGAATCTTGAG-3'	2430 2431 2432 2433
A456 Probe Set P i	FRET/TARGET SET 10 5'-CAGTCACGTCTTCAGGGAGTAGCGCA-NH2-3' 5'-CCCGTGGTAGGAGCAGCACTA-3' 3'-NH2- GCAGAGAGTCCCTCATCGCGT -5'	2434 2435 2436
C759 Probe Set p i a	FRET/TARGET SET 2 5'-CTCTCTGGTCTGGCCCACGGATT-NH2 5'-CTCCCACCAGTCGCTCACGTAATTTGTAA-3' 5'-AATCCTGGTGGGCGAGACG-B-3' 5'-TTAACTTCAAATACCACTGTAATCTTGGTCCAAGACCG-3'	2437 2438 2439 2440
G329 Probe Set p i	FRET/TARGET SET 4 5'-ACCGAGGCGCACCAATTATTCCTAACG-b-3' 5'-GCCGTTTCCAGAGTCCGATTGATTTTTGA-3' 3'-(biotin)- GCGGTGGTTAATAAGGATTGC -5'	2441 2442 2443
C1763 Probe Set p i a	FRET/TARGET SET 9 5'-CATCTTCGCGGAGACATTTCTTGATGATTCCTT-3' 5'-AAAGGTGTCTGGGCTCGTGCT-3' 3'-(bioitn)- GCCTCTGTAAAGAACTACTAAGGAA -5'	2444 2445 2446
Phosphatidylinositol-3-phosphate p110_, human (h-Pl3Kp110_) G1045 Probe Set (FV Arm) P 5'-Az I 5'-Cl	9110_) FRET/TARGET SET 4 5'-AACGAGCGCACCAGTTTCCTCTGTG-NH2-3' 5'-GACCAGCCCTGACATGAACTTTTAC-3' 3'-NH2- CGCGTGGTCAAAGGAGACAC -5'	2447 2448 2449
C1521 Probe Set Pp	FRET/TARGET SET 2 5-CTCTCTCGTCTCGGGAGGGTAATAATAAGG-NH2-3' 5-GCTGCCTTTTCAATAATCTTATCGAAC-3' 3'NH2-AGCAGGCCCTCCCATTATTATTCC-5'	2450 2451 2452
C2667 Probe Set p i	FRET/TARGET SET 2 5'-CTCTCTCGTCCGTTGTATTCTTTAAGCCAG-NH2-3' 5'-CGGTCCAGGTCATCCCCAGAC-3'	2453 2454

FIGURE 47BN

CU.	3'NH2-AGCAGCAACATAAGAAATTCGGTC-5'	2455
G537 Probe Set p i a	FRET/TARGET SET 2 5'-CTCTCTCGTCTCCTCGTGGATATGTTTG-NH2-3' 5'-CTAAGTTTTCAGGGATGGATGGTTCATGC-3' 3'NH2- AGCAGAGGACCATACAAAC -5'	2456 · 2457 2458
T3192 Probe Set p i a	FRET/TARGET SET 2 5'-CTCTCTCGTCTCAACTGTGGGC-NH2-3' 5'-TTAAGATCTGTAGTCTTTCCGAAC-3' 3'NH2- AGCAGAGTTCACACCCG- 5'	2459 2460 2461
Cartilage-derived morphogenic protein 1, human (h-CDMP1) A831 Probe Set P F F F F F F F F F F F F F F F F F F	11) FRET/TARGET SET 6 5'-CCGTCACGCCTCCTCCC-(biotin)-3' 5'-AGCCTCCAACTTCACGCTGT-3' 5'-AGCCTCCAACTCACGCTGT-3' 5'-GGGAGGCAACAGGAGGG-(biotin)-3'	2462 2463 2464
A1691 Probe Set p I a	FRET/TARGET SET 5 S'-CCGCCGAGATCACTGATGG-(biotin)-3' S'-ACACCACGTTGTTGGCAGAGTCAAG-3' S'-CCATCACCTTTTGCCAGAGTCAAG-3'	2465 2466 2467
b-actin, rat (r-bACT) C1671 Probe Set (longer) p i a	FRET/TARGET SET 6 5'-CCGTCACGCCTCGCCTTAGGGTTCA-NH2-3' 5'-TCTGGGTCATCTTTCACGGTTGA-3' 3'- GCGGAGCGCAATCCCAAGT -5' 5'-GAGGGGCCTCGGTGAGC-3'	2468 2469 2470 2471
Bile Salt port Pump, rat (r-BSEP) p p I	FRET/TARGET SET 5 5'-CCGCCGAGATCACGAGTTCTTGCCTTTC-(biotin)-3' 6'-CCGCCGAGATCACGAGTTCTTGCCTTTC-NH3-3' 5'-TTCACACACGCTTTCCTGGTATCTCC-3' 3'-(biotin)-CTAGTGCTCAAGAACGAAAG-5'	2472 2473 2474 2475
G1288 Probe Set p I a	FRET/TARGET SET 2 5'-CTCTCTCGTCTCCCAGAAGGCCAGT-(biotin)-3' 5'-TTCTTCATCTAGGACAAGTGTGGAACCATAA-3' 5'-ACTGGCCTTCTGGGAGACG-(biotin)-3'	2476 2477 2478

FIGURE 47BO

A790 Probe Set · · · · · · · · · · · · · · · · · · ·	FRET/TARGET SET 6 5'-CCGTCACGCCTCTTTCCTCCT-(biotin)-3' 5'-CCGAATTTCCATTCTCTCCGGAAGTAAATC-3' 5'-AGGAGAATGAGGAAGGGAAGTAAATC-3'	2479 2480 2481
Nitric Oxide Synthase 2A, human (h-iNOS2) A3418 Probe Set p I	FRET/TARGET SET 6 5'-CCGTCACGCCTCTGTCTTCTTCGC-(biotin)-3' 5'-GCTGCACCGCCACCCC-3' 5'- GCGAAGAAGACGACGCG-(biotin)-3'	2482 2483 2484
Neutral Carboxy Ester Hydrolase, human (h-NCEH) A1221 Probe Set p p i	FRET/TARGET SET 7 5:-AACGAGGCGCACTCTTCTTATTCTCCTG-B-3' 5:-AACGAGGCGCACTCTTCTTATTCTCCTG-NH2-3' 5:-GTCTCAAAGTCCACCAGTCTC-3' 5:-CAGGAGAATAAGAAGAGTGCGC-(biotin)-3'	2485 2486 2487 2488
A1221 Probe Set p i a	FRET/TARGET SET 6 5'-CCGTCACGCCTCTTCTTATTCTCC-3' 5'-CCGTCACGCCTCTTCTTATTCTCC-NH2-3' 5'-CTCTCAAGTCCACCAGTCTC-3' 3'-GCGAGAGAATAAGAGG-5' 5'-TGGGATGGGTCCTGGGC-3'	2489 2490 2491 2492 2493
C1309. Probe Set P i a	FRET/TARGET SET 8 5'-GAACGGCAGGTTTGGCACTCTTGGCATT-NH2-3' 5'- CAGGTAGGCG TAGGTCTTGA-3' 3'-NH2-CGTCCAAACCGTGAGAACCGTAA-5' 5'-GGCTCTGTGCTGGGCTA-NH2-3'	2494 2495 2496 2497
Peroxisomal Proliferation Activator Protein Receptor alpha, human (h-PPAR_) G1480 Probe Set FRET/TARGET SE p 5'-CGTCACGCC p 5'-CGGTCACGCC l 5'-CGGGTGCAGC a 5'-AGACGCAGT a	, human (h-PPAR_) FRETJTARGET SET 6 5'-CCGTCACGCCTCCCGACTCCGTCT-(biotin)-3' 5'-CGGGTGCAGCGCAGCATT-3' 5'-AGACGGAGTCGGGAGGCG-(biotin)-3'	2498 2499 2500
A1044 Probe Set p i a	FRET/TARGET SET 6 5'-CCGTCACGCCTCTGTCACTTGTTCT-(biotin)-3' 5'-TGGCCTCATAAACTCCGTATTTTAGCAAG-3' 5'-AGAACGATCAAGTGACAGAGCG-(biotin)-3'	2501 2502 2503

FIGURE 47BP

C 1311 Probe Set p i	FRET/TARGET SET 6 5'-CCGCCGAGATCACGTTTAGAAG-(biotin)-3' 5'-CACATGTACAATACCTCCTGCATTTTTTCAATC-3' 5'-CTTCTAAACGTAGGACACGTGATCTCGG-(biotin)-3'	2504 2505 2506
Peroxisomal Proliferation Activator Protein Receptor beta, human (h-PPAR_A595 Probe set 6B. Designed truncated probe and stackers to reduce temperature p 5-CCGTCACGC i 5-CTGGCACTT a 3-NH2-GCGGC s	human (h-PPAR_) FRET/TARGET SET 6 Serature S-CCGTCACGCCTCTTCTGAATCTTGC-3' 5'-CTGGCACTTGTTGCGGTTCTA-3' 3'-NH2-GCGGAGAAGACTTAGAACG-5' 5'-AGCTGCGCTCACACTTCTCGT-3'	2507 2508 2509 2510
6C Decim for new INIVADED accommittee 609, 21 Ma	FRET/TARGET SET 6	
	5'-CCGTCACGCCTCTTCTGAATCTTG-NH2-3' 5'- CTGGCACTTGT TGCGGTTCTA-3' 3'-NH2- GCGGAGAAGACTTAGAAC -5' 5'-CAGCTGCGCTCACACTTCTCGT-NH2-3'	2511 2512 2513 2514
6D. Truncate probe. p i s	FRET/TARGET SET 6 5'-CCGTCACGCCTCTTCTGAATCTT-NH2-3' 5'-CCTGGCACTTGTTGCGGTTCTA-3' 5'-CCTGGCACTTGTTGCGGTTCTA-3' 5'-GCAGCTGCGCTCACACTTCTCGT-NH2-3'	2515 2516 2517
C891 Probe Set p i a s	FRET/TARGET SET 7 5'-AACGAGGCGCACGGTAGGCATTGTAGA-3' 5'-CCTTCTTTTTGGTCATGTTGAAGTTTTTCAC-3' 3'-CGCGTGCCATCCGTAACATCT-5' 5'-TGTGCTGGAGGAGCTTCA-3'	2518 2519 2520 2521
Substance P, rat (r-SubP) C344 Probe Set P I a	FRET/TARGET SET 6 5'-CCGTCACGCCTCGCCACTTGTTTTTCA-NH2-3' 5'-CCATGCCCATAAAGAGCCTTTAACAGGA-3' 3'-NH2- GCGGAGCGGTGAACAAAAGT -5' NO STACKER	2522 2523 2524
A396 Probe Set P	FRET/TARGET SET 6 5'-CCGTCACGCCTCTTTATGCCTTTTGTGA-NH2-3'	2525

FIGURE 47BQ

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@ v	5'-TGCCCATTAGTCCAACAAAGGAATCTGTA-3' 3'-GCGGAGAATACGGAAACACT-5' 5'-GAGATCTGACCATGCCCATAAAGAGCC-NH2-3'	2526 2527 2528
C752 Probe Set p . i	FRET/TARGET SET 7 5'-AACGAGGCGCACGCTGGCAACTTGT-NH2-3' 5'-CCTTTCTGTCTTTGGAGACTTGCATCA-3' 3'-NH2-CGCGTGCGACCGTTTGAACA-5' 5'-ACAACTCCATCAACACTTGCTG-NH2-3'	2529 2530 2531 2532
Hepatic Lipase, human (h-LIPC) A830 Probe Set p i i a	FRET/TARGET SET 7 5-AACGAGGCGCACTCTAGGAAGTGGCA-NH2-3' 5-GTGCTGGGCAATATGTCTGTAGAGCG-3' 3'-NH2-CGCGTGAGATCCTTCACCGT-5' 5'-GCCAGGCTGGAAGGAGC-NH2-3'	2533 2534 2535 2536
C1154 Probe Set p i a	FRET/TARGET SET 5 5'-CCGCCGAGATCACCGTCTCAGTTTGGT-NH2-3' 5'-CGAGTAGTGACATGGTAAAAGTTGTTTGTATTGGCT-3' 3'-NH2-CTCTAGTGCAGAGAGTCAAACCA-5'	2537 2538 2539
Hepatic Lipase, rat (r-LIPC) G357 Probe Set p i a	FRET/TARGET SET 5 5'-CCGCCGACATCACCGGGTT-NH2-3' 5'-GGGAGATCCACTCCACGGGTT-NH2-3' 3'-NH2-TCTAGTGGTGCCAGTGCCCAA-5' 5'-GGGACTGTCGGACTTCAGG-NH2-3'	2540 2541 2542 2543
C1167 Probe Set p i a s	FRET/TARGET SET 8 5'-GAACGGCAGGTTTGGGGGAATTTTCTTTATTTCTT-NH2-3' 5'-ATTCCTTCGCCCAGGGTGATG-3' 3'-NH2-GTCCAAACCCCTTAAAAGAAATAAAGAA-5' 5'-CTTTTGTCCCCAGCGGTGT-NH2-3'	2544 2545 2546 2546
Metabotropic Glutamate Receptor 2, rat (r-mGluR2) C1403 Probe Set P I a	FRET/TARGET SET 7 5'-AACGAGGCGCACGGTGGTGTTGGGA-NH2-3' 5'-GCCTCATAGCATCGCAGAGGTGT-3' 3'-NH2-CGCGTGCCACCACACCT-5' 5'-CAGAGGGCACGTGTTGT-NH2-3'	2548 2549 2550 2551

FIGURE 47BR

G-protein coupled receptor 2, rat (r-ETBR-LP2) A1629 Probe set p I a	FRET/TARGET SET 8 5'-GAACGGCAGGTTTGTCAGCAGACCGC-NH2-3' 5'-GAGAGGCCAAAGTGAGACCATGTGAAAGAAA-3' 3'-NH2-CGTCCAAACAGTCGTCTGGCG-5' 5'-CATGGATCGGCATGTCGCTCTGGCG-5'	2552 2553 2554 2555
i kappa b alpha, human (h-MAD3) C542 Probe Set p I	FRET/TARGET SET 7 5'-AACGAGGCGCACGGTGTAGGGGGG-(biotin)-3' 5'-GCCCTGCTCACAGGCAAT-3' 5'-CCCCCTACACGTGCGC-(biotin)-3'	2556 2557 2558
C363 Probe Set P I	FRET/TARGET SET 6 5-CCGTCACGCCTCGTCAGTGCCTTTTC-(biotin)-3' 5-CACCTGGCGGATCACTTCCATGT 5-GAAAAGGCACTGACGAGGCG-(biotin)-3'	2559 2560 2561
G953 Probe Set P I A	FRET/TARGET SET 6 5'-CCGTCACCCTCCTCACT-(biotin)-3' 5'-ACTCTGACTCTGTCATAGCTCTT 5'-AGTGAGGATGAGGGGGG(biotin)-3'	2562 2563 2564
C923 Probe Set P I A A S	FRET/TARGET SET 7 5'-AACGAGGCGCACGGTTTTCTAGTGTCA-NH2-3' 5'-CTCACTCTGGCAGCATCTGAAT-3' 3'-NH2-CGCGTGCCAAAGATCACAGT-5' 5'-GCTGGCCAGCGCONH2-3'	2565 2566 2567 2568
Lecithin cholesterol acyltransferase, human (h-LCAT) C821 Probe Set (truncated Probe Design) p l a s	FRET/TARGET SET 5 5'-CCGCCGAGATCACGGTTATGCGCTG-NH2-3' 5'-CCAGGGGGAGGTGGTC-3' 3'-NH2-TCTAGTGCCAATACGCGACG-5' 5'-CTCCTTTTCAGTGCTGGTGG-NH2-3'	2569 2570 2571 2572
C827 Probe Design p I a	FRET/TARGET SET 8 5'-GAACGGCAGGTTTGGGTGGTTATGCG-NH2-3' 5'-AGAGGAAACATCCAGGGGGAG-3' 3'-NH2-CGTCCAAACCACCACCAATACGC-5'	2573 2574 2575

FIGURE 47BS

C1217 Probe Design p I	FRET/TARGET SET 5 5'-CCGCCGAGATCACGAGATGCTGTATCCC-NH2-3' 5'-GGTCAGGTTGCTGAAGACCATGTTG-3' 3'-NH2-TCTAGTGCTCTACGACATAGGG-5'	2576 2577 2578
Apolipoprotein A-1, human (h-ApoA1) A177 Probe Set p l a s	FRET/TARGET SET 6 5'-CCGTCACGCCTCTGAGCACATCCACG-NH2-3' 5'-ACATGTCTCTGCCGCTGTCTTA-3' 3'-NH2-GCGAGACTCGTGTAGGTGC-5' 5'-TACACAGTGGCCAGGTCCTT-NH2-3'	2579 2580 2581 2582
A227 Probe Set (titrate length of 2'-O-Me in Invader) p i i A s	FRET/TARGET SET 8 5'-GAACGGCAGGTTTGTCCCAAGGCGG-NH2-3' 5'-GTCAAGGACTTTAGGTTTAGCTGTTTA-3' 5'-GTCAAGGATCTTTAGGTTTAGCTGTTTA-3' 5'-GTCCAGGTTCTTTAGGTTTAGGTTTAGCTTTTA-3' 5'-GTCCCAGTTGTCAAGGATCTTTAGGTTTAGCTTTTA-3' 3'-NH2-GTCCAAACAGGGTTCCGCC-5' 5'-AGCCTTCAAACTGGGACACATAGTCTC-NH2-3'	2583 2584 2585 2586 2587 2588
G350 Probe Set P I a	FRET/TARGET SET 5 5'-CCGCCGAGATCACTTCTGTCTCCTT-NH2-3' 5'-CTCCTGCCTCAGGCCG-3' 3'-NH2 -TCTAGTGGAGAGGAA -5' 5'- TTCCAGGTTATCCCAGAACTCC -NH2-3'	2589 2590 2591 2592
G233 Probe Set p I a s	FRET/TARGET SET 11 5'-AGAACGGCAGTCTTTCCCAAGG-NH2-3' 5'-CCAGTTGTCAAGGCTTTAGGTTTAGT-3' 3'-NH2-CGTCAGAAAGGCTTAGGTTCC-5' 5'-CGGAGCCTTCAAAGGGTTCC-5'	2593 2594 2595 2596
Metabotropic Glutamate Receptor 1, rat (r-mGluR1) T934 Probe Set p I	FRET/TARGET SET 11 5'-AGAACGGCAGTCTTTAGAATAGGCGATCTGT-NH2-3' 5'-CACTCAGGTCTATGCTGTGGCT-3' 3'-NH2-GTCAGAATCTTATCCGCTAGACA-5' 5'-GGGATGTCGAACAGAGATTCT-NH2-3'	2597 2598 2599 2600

Ubiquitin, human (h-UBIQ)

FIGURE 47BT

G119 Probe Set (MO4 Arm) p I a	FRET/TARGET SET 6 5'-CCGTCACGCCTCCTTTACATTTCTATCGTATCCG-(biotin)-3' 5'-CCTTCCTTATCCTGGATCTTGGCA-3' 3'-(biotin)-GCGGAGGAAATGTAAAAGATAGCATAGGC-5'	2601 2602 2603
G119 Probe Set p i a	FRET/TARGET SET 5 5'-CGCCGAGATCACCTTTACATTTCTATCGTATCCG-(biotin)-3' 5'-CCTTCCTTATCCTGGATCTTGGCA-3' 3'-(biotin)-CTAGTGGAAATGTAAAAGATAGCATAGGC-5'	2604 2605 2606
G131 Probe Set p I a	FRET/TARGET SET 9 5'-CATCTTCGCGGACTGGATCTTGGCC-(biotin)-3' 5'-GCTGATCAGGAGGAATTCCTTCCTTATCT-3' 3'-(biotin)-GCCTGACCTAGAACCGG-5'	2607 2608 2609
Scanned G119 region (ELISA format (No Arrestors) p p p l	5'-CTCTCTCGTCTTTACATTTTCTATCGTATCCGA-NH2-3' 5'-CTCTCTCGTCTTTACATTTTCTATCGTATCCGA-NH2-3' 5'-CTCTCTCGTCTCTTTACATTTTCTATCGTATCCG-NH2-3' 5'-CTCTCTCGTCTCCCTTTACATTTTCTATCGTATC-NH2-3' 5'-CTCTCTCGTCTCCCTTTACATTTTCTATCG-NH2-3' 5'-GGAATTCCTTCCTTACATTTTCTATCG-NH2-3' 5'-GGAATTCCTTCCTTATCCTGGATCTTGGC-3' 5'-CCTTCCTTATCCTGGATCTTGGCA-3' 5'-TCCTTATCCTGGATCTTGGCCA-3' 5'-TCCTTATCCTGGATCTTGGCCA-3'	2610 2611 2612 2613 2614 2615 2616 2618
Ubiquitin, mouse (m-UBIQ) G294 Probe Set p I	FRET/TARGET SET 7 5'-CCGTCACGCCTCCCTTCTGGATGTTGTA-(biotin)-3' 5'-CCAGGTGCAGGGTTGACTA-3' 3'-(biotin)-GCGGAGGGAAGACCTA-5'	2620 2621 2622
G294 Probe Set p I a	FRET/TARGET SET 5 5'-CGCCGAGATCACCCTTCTGGATGTTGTA-(biotin)-3' 5'-CCAGGTGCAGGGTTGACTA-3' 3'-(biotin)-CTAGTGGGAAGACCTACAACAT-5'	2623 2624 2625
G294 Probe Set p I	FRET/TARGET SET 6 5'-CCGTCACGCCTCCCTTCTGGATGTTGTAAT-NH2-3' 5'-CCAGGTGCAGGGTTGACTA-3'	2626 2627

FIGURE 47BU

' ro	3'-NH2-GCGGAGGGAAGACCTACAACATTA-5'	2628
G294 Probe Set p i a	FRET/TARGET SET 6 5'-CCGTCACGCCTCCCTTCTGGATGTTGTAATC-NH2-3' 5'-CCAGGTGCAGGGTTGACTA-3' 3'-NH2- GCGAGGGAAGACCTACAACATTAG -3'	2629 2630 2631
T514 Probe Set p i a	FRET/TARGET SET 7 5'-AACGAGGCGCACATGTTGTAATCAGAGAGGG-NH2-3' 5'-TGCAGGGTTGACTCTTTCTGGA-3' 3'-NH2-CGCGTGTACAACATTAGTCTTCCCC-5'	2632 2633 2634
G750 Probe Set p	FRET/TARGET SET 9 5'-CATCTTCGGGACCTTCTGGATGTTGTA-NH2-3' 5'-GGACCAGGTGCAGGGTTGACTT-3' 3'-NH2-GCCTGGAAGACCTACAACAT-5'	2635 2636 2637
G185 Probe Set p I a	FRET/TARGET SET 9 5'-CATCTTCGCGGACTTCACGTTCTCGATGG-NH2-3' 5'-CCCTCTTTATCCTGGATCTTGGCA-3' 3'-NH2-GCGCCTGAAGTGCAAGAGCTACC-5'	2638 2639 2640